## SEQUENCE LISTING

<110> Genentech, Inc. Ashkenazi, Avi Botstein, David Desnoyers, Luc Eaton, Dan L Ferrara, Napoleone Filvaroff, Ellen Fong, Sherman Gao, Wei-Qiang Gerber, Hanspeter Gerritsen, Mary\E. Goddard, A. Godowski, Paul J Grimaldi, Christopher J. Gurney, Austin L.\ Hillan, Kenneth, Kljavin, Ivar J. Mather, Jennie P. Pan, James Paoni, Nicholas F. Roy, Margaret Ann Stewart, Timothy A. Tumas, Daniel Williams, P. Mickey Wood, William, I.

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<213> Homo sapiens
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<400> 18

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Pro Pro Ser Leu Arg Cys Ser Leu His Ser Ala Cys Cys Ser Gly Asp

Pro Ala Ser Tyr Arg Leu Trp Gly Ala Pro Leu Gln Pro Thr Leu Gly 55

Val Val Pro Gln Ala Ser Val Pro Leu Leu Thr Asp Leu Ala Gln Trp 70

Glu Pro Val Leu Val Pro Glu Ala His Pro Asn Ala Ser Leu Thr Met

Tyr Val Cys Thr Pro Val Pro His Pro Asp Pro Pro Met Ala Leu Ser 100 105

Arg Thr Pro Thr Arg Gln Ile Ser Ser Ser Asp Thr Asp Pro Pro Ala 120

Asp Gly Pro Ser Asn Pro Leu Cys Cys Cys Phe His Gly Pro Ala Phe 135 130

Ser Thr Leu Asn Pro Val Leu Arg His Leu Phe Pro Gln Glu Ala Phe 150

Pro Ala His Pro Ile Tyr Asp Leu Ser Gln Val Trp Ser Val Val Ser 170 165

Pro Ala Pro Ser Arg Gly Gln Ala Leu Arg Arg Ala Gln 180

<210> 19

<211> 24

<212> DNA

<213> Artificial Sequence

<223> Description of Artificial Sequence: Synthetic oligonucleotide probe

<400> 19

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<210> 20

<211> 24

<212> DNA

24

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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<212> PRT
<213> Homo sapiens
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<400> 23

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Arg Thr Ser Gly Lys His Val Gln Val Thr Gly Arg Arg Ile Ser Ala
50 55 60

Thr Ala Glu Asp Gly Asn Lys Phe Ala Lys Leu Ile Val Glu Thr Asp 65 70 75 80

Thr Phe Gly Ser Arg Val Arg Ile Lys Gly Ala Glu Ser Glu Lys Tyr 85 90 95

Ile Cys Met Asn Lys Arg Gly Lys Leu Ile Gly Lys Pro Ser Gly Lys
100 105 110

Ser Lys Asp Cys Val Phe Thr Glu Ile Val Leu Glu Asn Asn Tyr Thr 115 120 125

Ala Phe Gln Asn Ala Arg His Glu Gly Trp Phe Met Ala Phe Thr Arg 130 135 140

Gln Gly Arg Pro Arg Gln Ala Ser Arg Ser Arg Gln Asn Gln Arg Glu 145 150 155 160

Ala His Phe Ile Lys Arg Leu Tyr Gln Gly Gln Leu Pro Phe Pro Asn 165 170 175

His Ala Glu Lys Gln Lys Gln Phe Glu Phe Val Gly Ser Ala Pro Thr 180 185 190

Arg Arg Thr Lys Arg Thr Arg Arg Pro Gln Pro Leu Thr 195 200 205

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<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic oligonucleotide probe

<400> 24

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<211> 24
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 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Synthetic
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<210> 26
<211> 41
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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<222> (21)
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<212> DNA
<213> Homo sapiens
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Leu Leu Ala Cys Pro Ser Val Cys Arg Cys Asp Arg Asn Phe Val Tyr
                             40
Cys Asn Glu Arg Ser Leu Thr Ser Val Pro Leu Gly Ile Pro Glu Gly
     50
                         55
Val Thr Val Leu Tyr Leu His Asn Asn Gln Ile Asn Asn Ala Gly Phe
                     70
                                         75
Pro Ala Glu Leu His Asn Val Gln Ser Val His Thr Val Tyr Leu Tyr
                 85
Gly Asn Gln Leu Asp Glu Phe Pro Met Asn Leu Pro Lys Asn Val Arg
                                105
Val Leu His Leu Gln Glu Asn Asn Ile Gln Thr Ile Ser Arg Ala Ala
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120

2479

- Leu Ala Gln Leu Leu Lys Leu Glu Glu Leu His Leu Asp Asp Asn Ser 130 135 140
- Leu Lys Leu Phe Leu Ser Lys Asn His Leu Ser Ser Val Pro Val
  165 170 175
- Gly Leu Pro Val Asp Leu Gln Glu Leu Arg Val Asp Glu Asn Arg Ile 180 185 190
- Ala Val Ile Ser Asp Met Ala Phe Gln Asn Leu Thr Ser Leu Glu Arg 195 200 205
- Leu Ile Val Asp Gly Asn Leu Leu Thr Asn Lys Gly Ile Ala Glu Gly 210 215 220
- Thr Phe Ser His Leu Thr Lys Leu Lys Glu Phe Ser Ile Val Arg Asn 225 230 235 240
- Ser Leu Ser His Pro Pro Pro Asp Leu Pro Gly Thr His Leu Ile Arg 245 250 255
- Leu Tyr Leu Gln Asp Asn Gln Ile Asn His Ile Pro Leu Thr Ala Phe 260 265 270
- Ser Asn Leu Arg Lys Leu Glu Arg Leu Asp Ile Ser Asn Asn Gln Leu 275 280 285
- Arg Met Leu Thr Gln Gly Val Phe Asp Asn Leu Ser Asn Leu Lys Gln 290 295 300
- Leu Thr Ala Arg Asn Asn Pro Trp Phe Cys Asp Cys Ser Ile Lys Trp 305 310 315 320
- Val Thr Glu Trp Leu Lys Tyr Ile Pro Ser Ser Leu Asn Val Arg Gly
  325 330 335
- Phe Met Cys Gln Gly Pro Glu Gln Val Arg Gly Met Ala Val Arg Glu 340 345 350
- Leu Asn Met Asn Leu Leu Ser Cys Pro Thr Thr Thr Pro Gly Leu Pro 355 360 365
- Leu Phe Thr Pro Ala Pro Ser Thr Ala Ser Pro Thr Thr Gln Pro Pro 370 375 380
- Thr Leu Ser Ile Pro Asn Pro Ser Arg Ser Tyr Thr Pro Pro Thr Pro 385 390 395 400
- Thr Thr Ser Lys Leu Pro Thr Ile Pro Asp Trp Asp Gly Arg Glu Arg

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|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val        | Thr        | Pro        | Pro<br>420 | Ile        | Ser        | Glu        | Arg        | Ile<br>425 | Gln        | Leu        | Ser        | Ile        | His<br>430 | Phe        | Val        |
| Asn        | Asp        | Thr<br>435 | Ser        | Ile        | Gln        | Val        | Ser<br>440 | Trp        | Leu        | Ser        | Leu        | Phe<br>445 | Thr        | Val        | Met        |
| Ala        | Tyr<br>450 | Lys        | Leu        | Thr        | Trp        | Val<br>455 | Lys        | Met        | Gly        | His        | Ser<br>460 | Leu        | Val        | Gly        | Gly        |
| Ile<br>465 | Val        | Gln        | Glu        | Arg        | Ile<br>470 | Val        | Ser        | Gly        | Glu        | Lys<br>475 | Gln        | His        | Leu        | Ser        | Leu<br>480 |
| Val        | Asn        | Leu        | Glu        | Pro<br>485 | Arg        | Ser        | Thr        | Tyr        | Arg<br>490 | Ile        | Cys        | Leu        | Val        | Pro<br>495 | Leu        |
| Asp        | Ala        | Phe        | Asn<br>500 | Tyr        | Arg        | Ala        | Val        | Glu<br>505 | Asp        | Thr        | Ile        | Суѕ        | Ser<br>510 | Glu        | Ala        |
| Thr        | Thr        | His<br>515 | Ala        | Ser        | Tyr        | Leu        | Asn<br>520 | Asn        | Gly        | Ser        | Asn        | Thr<br>525 | Ala        | Ser        | Ser        |
| His        | Glu<br>530 | Gln        | Thr        | Thr        | Ser        | His<br>535 | Ser        | Met        | Gly        | Ser        | Pro<br>540 | Phe        | Leu        | Leu        | Ala        |
| Gly<br>545 | Leu        | Ile        | Gly        | Gly        | Ala<br>550 | Val        | Ile        | Phe        | Val        | Leu<br>555 | Val        | Val        | Leu        | Leu        | Ser<br>560 |
| Val        | Phe        | Cys        | Trp        | His<br>565 | Met        | His        | Lys        | Lys        | Gly<br>570 | Arg        | Tyr        | Thr        | Ser        | Gln<br>575 | Lys        |
| Trp        | Lys        | Tyr        | Asn<br>580 | Arg        | Gly        | Arg        | Arg        | Lys<br>585 | Asp        | Asp        | Tyr        | Cys        | Glu<br>590 | Ala        | Gly        |
| Thr        | Lys        | Lys<br>595 | Asp        | Asn        | Ser        | Ile        | Leu<br>600 | Glu        | Met        | Thr        | Glu        | Thr<br>605 | Ser        | Phe        | Gln        |
| Ile        | Val<br>610 | Ser        | Leu        | Asn        | Asn        |            |            | Leu        |            |            | Gly<br>620 | Asp        | Phe        | Arg        | Leu        |
| Gln<br>625 | Pro        | Ile        | Tyr        | Thr        | Pro<br>630 | Asn        | Gly        | Gly        | Ile        | Asn<br>635 | Tyr        | Thr        | Asp        | Cys        | His<br>640 |
| Ile        | Pro        | Asn        | Asn        | Met<br>645 | Arg        | Tyr        | Cys        | Asn        | Ser<br>650 | Ser        | Val        | Pro        | Asp        | Leu<br>655 | Glu        |
| His        | Cys        | His        | Thr<br>660 |            |            |            |            |            |            |            |            |            |            |            |            |
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| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe   |           |
| <400> 29<br>cggtctacct gtatggcaac c   | 21        |
| <210> 30<br><211> 22<br><212> DNA<br><213> Artificial Sequence  |           |
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| <400> 30<br>gcaggacaac cagataaacc ac  | 22        |
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| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe   |           |
| <400> 31 acgcagattt gagaaggctg tc   | 22        |
| <210> 32 <211> 46 <212> DNA <213> Artificial Sequence   |           |
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| <210> 33<br><211> 3449<br><212> DNA<br><213> Homo sapiens   |           |
| <400> 33 acttggagca ageggeggeg geggagaeag aggeagagge agaagetggg geteegteet egeeteeeae gagegateee egaggagage egeggeeete ggegaggega | 60<br>120 |

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<211> 915

<212> PRT

<213> Homo sapiens

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Ser Ser Cys Glu Asn Lys Arg Ala Asp Leu Val Phe Ile Ile Asp Ser 50 55 60

Ser Arg Ser Val Asn Thr His Asp Tyr Ala Lys Val Lys Glu Phe Ile 65 70 75 80

Val Asp Ile Leu Gln Phe Leu Asp Ile Gly Pro Asp Val Thr Arg Val 85 90 95

Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys 100 105 110

Thr Phe Lys Arg Lys Ser Glu Val Glu Arg Ala Val Lys Arg Met Arg 115 120 125

His Leu Ser Thr Gly Thr Met Thr Gly Leu Ala Ile Gln Tyr Ala Leu 130 135 140

Asn Ile Ala Phe Ser Glu Ala Glu Gly Ala Arg Pro Leu Arg Glu Asn 145 150 155 160

Val Pro Arg Val Ile Met Ile Val Thr Asp Gly Arg Pro Gln Asp Ser 165 170 175

Val Ala Glu Val Ala Ala Lys Ala Arg Asp Thr Gly Ile Leu Ile Phe 180 185 190

Ala Ile Gly Val Gly Gln Val Asp Phe Asn Thr Leu Lys Ser Ile Gly
195 200 205

Ser Glu Pro His Glu Asp His Val Phe Leu Val Ala Asn Phe Ser Gln 210 215 220

Ile Glu Thr Leu Thr Ser Val Phe Gln Lys Lys Leu Cys Thr Ala His

| 225        |            |            |                    |            | 230        |            |            |            |            | 235        |            |            |            |            | 240        |
|------------|------------|------------|--------------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Met        | Cys        | Ser        | Thr                | Leu<br>245 | Glu        | His        | Asn        | Cys        | Ala<br>250 | His        | Phe        | Cys        | Ile        | Asn<br>255 | Ile        |
| Pro        | Gly        | Ser        | <b>T</b> yr<br>260 | Val        | Cys        | Arg        | Cys        | Lys<br>265 | Gln        | Gly        | Tyr        | Ile        | Leu<br>270 | Asn        | Ser        |
| Asp        | Gln        | Thr<br>275 | Thr                | Cys        | Arg        | Ile        | Gln<br>280 | Asp        | Leu        | Cys        | Ala        | Met<br>285 | Glu        | Asp        | His        |
| Asn        | Cys<br>290 | Glu        | Gln                | Leu        | Cys        | Val<br>295 | Asn        | Val        | Pro        | Gly        | Ser<br>300 | Phe        | Val        | Cys        | Gln        |
| Cys<br>305 | Tyr        | Ser        | Gly                | Tyr        | Ala<br>310 | Leu        | Ala        | Glu        | Asp        | Gly<br>315 | Lys        | Arg        | Cys        | Val        | Ala<br>320 |
| Val        | Asp        | Tyr        | Cys                | Ala<br>325 | Ser        | Glu        | Asn        | His        | Gly<br>330 | Cys        | Glu        | His        | Glu        | Cys<br>335 | Val        |
| Asn        | Ala        | Asp        | Gly<br>340         | Ser        | Tyr        | Leu        | Cys        | Gln<br>345 | Cys        | His        | Glu        | Gly        | Phe<br>350 | Ala        | Leu        |
| Asn        | Pro        | Asp<br>355 | Glu                | Lys        | Thr        | Cys        | Thr<br>360 | Arg        | Ile        | Asn        | Tyr        | Cys<br>365 | Ala        | Leu        | Asn        |
| Lys        | Pro<br>370 | Gly        | Cys                | Glu        | His        | Glu<br>375 | Cys        | Val        | Asn        | Met        | Glu<br>380 | Glu        | Ser        | Tyr        | Tyr        |
| Cys<br>385 | Arg        | Cys        | His                | Arg        | Gly<br>390 | Tyr        | Thr        | Leu        | Asp        | Pro<br>395 | Asn        | Gly        | Lys        | Thr        | Cys<br>400 |
| Ser        | Arg        | Val        | Asp                | His<br>405 | Cys        | Ala        | Gln        | Gln        | Asp<br>410 | His        | Gly        | Сув        | Glu        | Gln<br>415 | Leu        |
| Cys        | Leu        | Asn        | Thr<br>420         | Glu        | Asp        | Ser        | Phe        | Val<br>425 | Cys        | Gln        | Cys        | Ser        | Glu<br>430 | Gly        | Phe        |
| Leu        | Ile        | Asn<br>435 | Glu                | Asp        | Leu        | Lys        | Thr<br>440 | Cys        | Ser        | Arg        | Val        | Asp<br>445 | Tyr        | Cys        | Leu        |
| Leu        | Ser<br>450 | Asp        | His                | Gly        | Суз        | Glu<br>455 | Tyr        | Ser        | Cys        | Val        | Asn<br>460 | Met        | Asp        | Arg        | Ser        |
| Phe<br>465 | Ala        | Cys        | Gln                | Cys        | Pro<br>470 | Glu        | Gly        | His        | Val        | Leu<br>475 | Arg        | Ser        | Asp        | Gly        | Lys<br>480 |
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| His        | Ser        | Cys        | Val<br>500         | Ser        | Ser        | Glu        | Asp        | Ser<br>505 | Phe        | Val        | Cys        | Gln        | Cys<br>510 | Phe        | Glu        |

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- Cys Gln Ala Ile Asp His Gly Cys Glu His Ile Cys Val Asn Ser Asp 530 535 540
- Asp Ser Tyr Thr Cys Glu Cys Leu Glu Gly Phe Arg Leu Ala Glu Asp 545 550 555 560
- Gly Lys Arg Cys Arg Arg Lys Asp Val Cys Lys Ser Thr His His Gly 565 570 575
- Cys Glu His Ile Cys Val Asn Asn Gly Asn Ser Tyr Ile Cys Lys Cys 580 585 590
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- Thr Glu Gly Pro Ile Asp Leu Val Phe Val Ile Asp Gly Ser Lys Ser 610 615 620
- Leu Gly Glu Glu Asn Phe Glu Val Val Lys Gln Phe Val Thr Gly Ile 625 630 635 640
- Ile Asp Ser Leu Thr Ile Ser Pro Lys Ala Ala Arg Val Gly Leu Leu 645 650 655
- Gln Tyr Ser Thr Gln Val His Thr Glu Phe Thr Leu Arg Asn Phe Asn 660 665 670
- Ser Ala Lys Asp Met Lys Lys Ala Val Ala His Met Lys Tyr Met Gly 675 680 685
- Lys Gly Ser Met Thr Gly Leu Ala Leu Lys His Met Phe Glu Arg Ser 690 695 700
- Phe Thr Gln Gly Glu Gly Ala Arg Pro Leu Ser Thr Arg Val Pro Arg 705 710 715 720
- Ala Ala Ile Val Phe Thr Asp Gly Arg Ala Gln Asp Asp Val Ser Glu 725 730 735
- Trp Ala Ser Lys Ala Lys Ala Asn Gly Ile Thr Met Tyr Ala Val Gly
  740 745 750
- Val Gly Lys Ala Ile Glu Glu Glu Leu Gln Glu Ile Ala Ser Glu Pro 755 760 765
- Thr Asn Lys His Leu Phe Tyr Ala Glu Asp Phe Ser Thr Met Asp Glu
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- Ile Ser Glu Lys Leu Lys Lys Gly Ile Cys Glu Ala Leu Glu Asp Ser 785 790 795 800

| Asp                  | Gly                              | Arg          | Gln        | Asp<br>805 | Ser        | Pro        | Ala        | Gly        | Glu<br>810 | Leu        | Pro        | Lys        | Thr        | Val<br>815 | Gln        |    |
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| Gln                  | Pro                              | Thr          | Glu<br>820 | Ser        | Glu        | Pro        | Val        | Thr<br>825 | Ile        | Asn        | Ile        | Gln        | Asp<br>830 | Leu        | Leu        |    |
| Ser                  | Cys                              | Ser<br>835   | Asn        | Phe        | Ala        | Val        | Gln<br>840 | His        | Arg        | Tyr        | Leu        | Phe<br>845 | Glu        | Glu        | Asp        |    |
| Asn                  | Leu<br>850                       | Leu          | Arg        | Ser        | Thr        | Gln<br>855 | Lys        | Leu        | Ser        | His        | Ser<br>860 | Thr        | Lys        | Pro        | Ser        |    |
| Gly<br>865           | Ser                              | Pro          | Leu        | Glu        | Glu<br>870 | Lys        | His        | Asp        | Gln        | Cys<br>875 | Lys        | Cys        | Glu        | Asn        | Leu<br>880 |    |
| Ile                  | Met                              | Phe          | Gln        | Asn<br>885 | Leu        | Ala        | Asn        | Glu        | Glu<br>890 | Val        | Arg        | Lys        | Leu        | Thr<br>895 | Gln        |    |
| Arg                  | Leu                              | Glu          | Glu<br>900 | Met        | Thr        | Gln        | Arg        | Met<br>905 | Glu        | Ala        | Leu        | Glu        | Asn<br>910 | Arg        | Leu        |    |
| Arg                  | Tyr                              | Arg<br>915   |            |            |            |            |            |            |            |            |            |            |            |            |            |    |
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| <220<br><223         | 3> De                            |              | _          | on of      |            |            | cial       | Seq        | uence      | e: Sy      | ynthe      | etic       |            |            |            |    |
|                      | )> 35<br>accct                   |              | ttgt       | gaata      | ac to      | CC         |            |            |            |            |            |            |            |            |            | 23 |
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|                      | )> 36                            |              | tctai      | taget      | it a       | ב          |            |            |            |            |            |            |            |            |            | 22 |
| <210<br><211<br><212 | )> 35<br>L> 45<br>2> Di          | 7<br>5<br>NA |            | l Sec      |            |            |            |            |            |            |            |            |            |            |            |    |

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| Leu        | His        | Leu<br>35  | Pro        | Ala        | Asn        | Arg        | Leu<br>40  | Gln        | Ala        | Val        | Glu        | Gly<br>45  | Gly        | Glu        | Val        |
| Val        | Leu<br>50  | Pro        | Ala        | Trp        | Tyr        | Thr<br>55  | Leu        | His        | Gly        | Glu        | Val<br>60  | Ser        | Ser        | Ser        | Gln        |
| Pro<br>65  | Trp        | Glu        | Val        | Pro        | Phe<br>70  | Val        | Met        | Trp        | Phe        | Phe<br>75  | Lys        | Gln        | Lys        | Glu        | Lys<br>80  |
| Glu        | Asp        | Gln        | Val        | Leu<br>85  | Ser        | Tyr        | Ile        | Asn        | Gly<br>90  | Val        | Thr        | Thr        | Ser        | Lys<br>95  | Pro        |
| Gly        | Val        | Ser        | Leu<br>100 | Val        | Tyr        | Ser        | Met        | Pro<br>105 | Ser        | Arg        | Asn        | Leu        | Ser<br>110 | Leu        | Arg        |
| Leu        | Glu        | Gly<br>115 | Leu        | Gln        | Glu        | Lys        | Asp<br>120 | Ser        | Gly        | Pro        | Tyr        | Ser<br>125 | Cys        | Ser        | Val        |
| Asn        | Val<br>130 | Gln        | Asp        | Lys        | Gln        | Gly<br>135 | Lys        | Ser        | Arg        | Gly        | His<br>140 | Ser        | Ile        | Lys        | Thr        |
| Leu<br>145 | Glu        | Leu        | Asn        | Val        | Leu<br>150 | Val        | Pro        | Pro        | Ala        | Pro<br>155 | Pro        | Ser        | Cys        | Arg        | Leu<br>160 |
| Gln        | Gly        | Val        | Pro        | His<br>165 | Val        | Gly        | Ala        | Asn        | Val<br>170 | Thr        | Leu        | Ser        | Cys        | Gln<br>175 | Ser        |
| Pro        | Arg        | Ser        | Lys<br>180 | Pro        | Ala        | Val        | Gln        | Tyr<br>185 | Gln        | Trp        | Asp        | Arg        | Gln<br>190 | Leu        | Pro        |
| Ser        | Phe        | Gln<br>195 | Thr        | Phe        | Phe        | Ala        | Pro<br>200 | Ala        | Leu        | Asp        | Val        | Ile<br>205 | Arg        | Gly        | Ser        |
| Leu        | Ser<br>210 | Leu        | Thr        | Asn        | Leu        | Ser<br>215 | Ser        | Ser        | Met        | Ala        | Gly<br>220 | Val        | Tyr        | Val        | Суя        |
| Lys<br>225 | Ala        | His        | Asn        | Glu        | Val<br>230 | Gly        | Thr        | Ala        | Gln        | Cys<br>235 | Asn        | Val        | Thr        | Leu        | Glu<br>240 |
| Val        | Ser        | Thr        | Gly        | Pro<br>245 | Gly        | Ala        | Ala        | Val        | Val<br>250 | Ala        | Gly        | Ala        | Val        | Val<br>255 | Gly        |
| Thr        | Leu        | Val        | Gly<br>260 | Leu        | Gly        | Leu        | Leu        | Ala<br>265 | Gly        | Leu        | Val        | Leu        | Leu<br>270 | Tyr        | His        |
| Arg        | Arg        | Gly<br>275 | Lys        | Ala        | Leu        | Glu        | Glu<br>280 | Pro        | Ala        | Asn        | Asp        | Ile<br>285 | Lys        | Glu        | Asp        |

| Ala Il<br>29    |                | Pro        | Arg        | Thr        | Leu<br>295          | Pro        | Trp        | Pro        | Lys        | Ser<br>300 | Ser        | Asp        | Thr        | Ile        |     |
|-----------------|----------------|------------|------------|------------|---------------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|-----|
| Ser Ly<br>305   | s Asn          | Gly        | Thr        | Leu<br>310 | Ser                 | Ser        | Val        | Thr        | Ser<br>315 | Ala        | Arg        | Ala        | Leu        | Arg<br>320 |     |
| Pro Pr          | o His          | Gly        | Pro<br>325 | Pro        | Arg                 | Pro        | Gly        | Ala<br>330 | Leu        | Thr        | Pro        | Thr        | Pro<br>335 | Ser        |     |
| Leu Se          | r Ser          | Gln<br>340 | Ala        | Leu        | Pro                 | Ser        | Pro<br>345 | Arg        | Leu        | Pro        | Thr        | Thr<br>350 | Asp        | Gly        |     |
| Ala Hi          | s Pro<br>355   |            | Pro        | Ile        | Ser                 | Pro<br>360 | Ile        | Pro        | Gly        | Gly        | Val<br>365 | Ser        | Ser        | Ser        |     |
| Gly Le          |                | Arg        | Met        | Gly        | Ala<br>3 <b>7</b> 5 | Val        | Pro        | Val        | Met        | Val<br>380 | Pro        | Ala        | Gln        | Ser        |     |
| Gln Al<br>385   | a Gly          | Ser        | Leu        | Val<br>390 |                     |            |            |            |            |            |            |            |            |            |     |
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| <223>           | Descr<br>oligo |            |            |            |                     |            | Seq        | uenc       | e: S       | ynth       | etic       |            |            |            |     |
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|                 | oligo          |            |            |            |                     |            | •          |            |            |            |            |            |            |            |     |

| <400> 42<br>ggccacagca tcaaaacctt agaactcaat gtactggttc ctccagctcc              | 50 |
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2822

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Cys Glu Ile Arg Asn Gly Ile Glu Ala Cys Tyr Cys Asn Met Gly Phe
Ser Gly Asn Gly Val Thr Ile Cys Glu Asp Asp Asn Glu Cys Gly Asn
     50
Leu Thr Gln Ser Cys Gly Glu Asn Ala Asn Cys Thr Asn Thr Glu Gly
Ser Tyr Tyr Cys Met Cys Val Pro Gly Phe Arg Ser Ser Ser Asn Gln
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Asp Arg Phe Ile Thr Asn Asp Gly Thr Val Cys Ile Glu Asn Val Asn
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Ala Asn Cys His Leu Asp Asn Val Cys Ile Ala Ala Asn Ile Asn Lys
                           120
Thr Leu Thr Lys Ile Arg Ser Ile Lys Glu Pro Val Ala Leu Leu Gln
                                           140
    130
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Glu Val Tyr Arg Asn Ser Val Thr Asp Leu Ser Pro Thr Asp Ile Ile
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                   150
Thr Tyr Ile Glu Ile Leu Ala Glu Ser Ser Ser Leu Leu Gly Tyr Lys
                                   170
                                                      175
                165
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Asn Asn Thr Ile Ser Ala Lys Asp Thr Leu Ser Asn Ser Thr Leu Thr

|            |            |            | 180        |            |            |            |            | 185        |            |            |            |            | 190        |            |            |
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| Glu        | Phe        | Val<br>195 | Lys        | Thr        | Val        | Asn        | Asn<br>200 | Phe        | Val        | Gln        | Arg        | Asp<br>205 | Thr        | Phe        | Val        |
| Val        | Trp<br>210 | Asp        | Lys        | Leu        | Ser        | Val<br>215 | Asn        | His        | Arg        | Arg        | Thr<br>220 | His        | Leu        | Thr        | Lys        |
| Leu<br>225 | Met        | His        | Thr        | Val        | Glu<br>230 | Gln        | Ala        | Thr        | Leu        | Arg<br>235 | Ile        | Ser        | Gln        | Ser        | Phe<br>240 |
| Gln        | Lys        | Thr        | Thr        | Glu<br>245 | Phe        | Asp        | Thr        | Asn        | Ser<br>250 | Thr        | Asp        | Ile        | Ala        | Leu<br>255 | Lys        |
| Val        | Phe        | Phe        | Phe<br>260 | Asp        | Ser        | Tyr        | Asn        | Met<br>265 | Lys        | His        | Ile        | His        | Pro<br>270 | His        | Met        |
| Asn        | Met        | Asp<br>275 | Gly        | Asp        | Tyr        | Ile        | Asn<br>280 | Ile        | Phe        | Pro        | Lys        | Arg<br>285 | Lys        | Ala        | Ala        |
| Tyr        | Asp<br>290 | Ser        | Asn        | Gly        | Asn        | Val<br>295 | Ala        | Val        | Ala        | Phe        | Leu<br>300 | Tyr        | Tyr        | Lys        | Ser        |
| Ile<br>305 | Gly        | Pro        | Leu        | Leu        | Ser<br>310 | Ser        | Ser        | Asp        | Asn        | Phe<br>315 | Leu        | Leu        | Lys        | Pro        | Gln<br>320 |
| Asn        | Tyr        | Asp        | Asn        | Ser<br>325 | Glu        | Glu        | Glu        | Glu        | Arg<br>330 | Val        | Ile        | Ser        | Ser        | Val<br>335 | Ile        |
| Ser        | Val        | Ser        | Met<br>340 | Ser        | Ser        | Asn        | Pro        | Pro<br>345 | Thr        | Leu        | Tyr        | Glu        | Leu<br>350 | Glu        | Lys        |
| Ile        | Thr        | Phe<br>355 | Thr        | Leu        | Ser        | His        | Arg<br>360 | Lys        | Val        | Thr        | Asp        | Arg<br>365 | Tyr        | Arg        | Ser        |
| Leu        | Cys<br>370 | Ala        | Phe        | Trp        | Asn        | Tyr<br>375 | Ser        | Pro        | Asp        | Thr        | Met<br>380 | Asn        | Gly        | Ser        | Trp        |
| Ser<br>385 | Ser        | Glu        | Gly        | Cys        | Glu<br>390 | Leu        | Thr        | Tyr        |            | Asn<br>395 |            | Thr        | His        | Thr        | Ser<br>400 |
| Cys        | Arg        | Cys        | Asn        | His<br>405 | Leu        | Thr        | His        | Phe        | Ala<br>410 | Ile        | Leu        | Met        | Ser        | Ser<br>415 | Gly        |
| Pro        | Ser        | Ile        | Gly<br>420 | Ile        | Lys        | Asp        | Tyr        | Asn<br>425 | Ile        | Leu        | Thr        | Arg        | Ile<br>430 | Thr        | Glr        |
| Leu        | Gly        | Ile<br>435 | Ile        | Ile        | Ser        | Leu        | Ile<br>440 | Cys        | Leu        | Ala        | Ile        | Cys<br>445 | Ile        | Phe        | Thr        |
| Phe        | Trp        |            | Phe        | Ser        | Glu        | Ile<br>455 |            | Ser        | Thr        | Arg        | Thr<br>460 | Thr        | Ile        | His        | Lys        |

Asn Leu Cys Cys Ser Leu Phe Leu Ala Glu Leu Val Phe Leu Val Gly
465 470 475 480

Ile Asn Thr Asn Thr Asn Lys Leu Phe Cys Ser Ile Ile Ala Gly Leu
485 490 495

Leu His Tyr Phe Phe Leu Ala Ala Phe Ala Trp Met Cys Ile Glu Gly 500 505 510

Ile His Leu Tyr Leu Ile Val Val Gly Val Ile Tyr Asn Lys Gly Phe 515 520 525

Leu His Lys Asn Phe Tyr Ile Phe Gly Tyr Leu Ser Pro Ala Val Val 530 535 540

Val Gly Phe Ser Ala Ala Leu Gly Tyr Arg Tyr Tyr Gly Thr Thr Lys 545 550 555 560

Val Cys Trp Leu Ser Thr Glu Asn Asn Phe Ile Trp Ser Phe Ile Gly 565 570 575

Pro Ala Cys Leu Ile Ile Leu Val Asn Leu Leu Ala Phe Gly Val Ile 580 585 590

Ile Tyr Lys Val Phe Arg His Thr Ala Gly Leu Lys Pro Glu Val Ser 595 600 605

Cys Phe Glu Asn Ile Arg Ser Cys Ala Arg Gly Ala Leu Ala Leu Leu 610 615 620

Phe Leu Leu Gly Thr Thr Trp Ile Phe Gly Val Leu His Val Val His 625 630 635

Ala Ser Val Val Thr Ala Tyr Leu Phe Thr Val Ser Asn Ala Phe Gln 645 650 655

Gly Met Phe Ile Phe Leu Phe Leu Cys Val Leu Ser Arg Lys Ile Gln 660 665 670

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Leu Arg 690

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aattetgaag aggaggaaag agteatatet teagtaattt eagteteaat gageteaaac 240
ccacccacat tatatgaact tgaaaaaata acatttacat taagtcatcg aaaggtcaca 300
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tggtcttcag agggctgtga gctgacatac tcaaatgaga cccacacctc atgccgctgt 420
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tataatatte ttacaaggat caeteaaeta ggaataatta tttcaetgat ttgtettgee 540
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| <210> 55 <211> 18 <212> DNA <213> Artificial Sequence  |    |
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| <210> 56 <211> 23 <212> DNA <213> Artificial Sequence  |    |
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cggcccggag ccgccgcgcc gtcagagcag gagcgctgcg tccaggatct agggccacga 360
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accccatcg ccggagctgc gccgagagcc ccagggaggt gccatgcgga gcgggtgtgt 480
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His Val His Tyr Gly Trp Gly Asp Pro Ile Arg Leu Arg His Leu Tyr 35

Thr Ser Gly Pro His Gly Leu Ser Ser Cys Phe Leu Arg Ile Arg Ala 60

| Asp<br>65  | Gly            | Val        | Val        | Asp          | Cys<br>70  | Ala        | Arg        | Gly        | Gln        | Ser<br>75  | Ala        | His        | Ser        | Leu        | Leu<br>80  |    |
|------------|----------------|------------|------------|--------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|----|
| Glu        | Ile            | Lys        | Ala        | Val<br>85    | Ala        | Leu        | Arg        | Thr        | Val<br>90  | Ala        | Ile        | Lys        | Gly        | Val<br>95  | His        |    |
| Ser        | Val            | Arg        | Tyr<br>100 | Leu          | Cys        | Met        | Gly        | Ala<br>105 | Asp        | Gly        | Lys        | Met        | Gln<br>110 | Gly        | Leu        |    |
| Leu        | Gln            | Tyr<br>115 | Ser        | Glu          | Glu        | Asp        | Cys<br>120 | Ala        | Phe        | Glu        | Glu        | Glu<br>125 | Ile        | Arg        | Pro        |    |
| Asp        | Gly<br>130     | Tyr        | Asn        | Val          | Tyr        | Arg<br>135 | Ser        | Glu        | Lys        | His        | Arg<br>140 | Leu        | Pro        | Val        | Ser        |    |
| Leu<br>145 | Ser            | Ser        | Ala        | Lys          | Gln<br>150 | Arg        | Gln        | Leu        | Tyr        | Lys<br>155 | Asn        | Arg        | Gly        | Phe        | Leu<br>160 |    |
| Pro        | Leu            | Ser        | His        | Phe<br>165   | Leu        | Pro        | Met        | Leu        | Pro<br>170 | Met        | Val        | Pro        | Glu        | Glu<br>175 | Pro        |    |
|            | Asp            |            | 180        |              |            |            |            | 185        |            |            |            |            | 190        |            |            |    |
| Glu        | Thr            | Asp<br>195 |            | Met          | Asp        | Pro        | Phe<br>200 | Gly        | Leu        | Val        | Thr        | Gly<br>205 | Leu        | Glu        | Ala        |    |
| Val        | Arg<br>210     |            | Pro        | Ser          | Phe        | Glu<br>215 |            |            |            |            |            |            |            |            |            |    |
|            | 0 > 6          |            |            |              |            |            |            |            |            |            |            |            |            |            |            |    |
|            | 1> 2<br>2> D   |            |            |              |            |            |            |            |            |            |            |            |            |            |            |    |
|            | .3> A          |            | icia       | l Se         | quen       | ce         |            |            |            |            |            |            |            |            |            |    |
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|            | 0> 6           |            | ataa       | ctac         | aa t       | atat       | a          |            |            |            |            |            |            |            |            | 26 |
| acc        | egec           | cay        | acgg       | ccac         | aa c       | 9090       | <b>~</b>   |            |            |            |            |            |            |            |            |    |
|            | .0> 6<br>.1> 4 |            |            |              |            |            |            |            |            |            |            |            |            |            |            |    |
| <23        | .2> D          | NA         |            |              |            |            |            |            |            |            |            |            |            |            |            |    |
| <23        | .3> A          | rtif       | icia       | .1 Se        | quen       | .ce        |            |            |            |            |            |            |            |            |            |    |
| <22<br><22 | 23 > D         |            |            | on o<br>eoti |            |            |            | . Seq      | uenc       | e: S       | ynth       | etic       | !          |            |            |    |
|            | 00> €          |            | _,         |              |            | ځيم        |            |            | ,,,,,,,,   | 12C+C      | r +=       |            |            |            |            | 42 |
| gc         | ctccc          | ggt        | CTCC       | ctga         | .gc a      | gugo       | cada       | ic ag      | regge      | ague       | Jua        |            |            |            |            |    |

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<210> 63
<211> 1295
<212> DNA
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- Ala Cys Lys Thr Pro Lys Lys Thr Val Ser Ser Arg Leu Glu Trp Lys
  50 55 60
- Lys Leu Gly Arg Ser Val Ser Phe Val Tyr Tyr Gln Gln Thr Leu Gln 65 70 75 80
- Gly Asp Phe Lys Asn Arg Ala Glu Met Ile Asp Phe Asn Ile Arg Ile 85 90 95
- Lys Asn Val Thr Arg Ser Asp Ala Gly Lys Tyr Arg Cys Glu Val Ser 100 105 110
- Ala Pro Ser Glu Gln Gly Gln Asn Leu Glu Glu Asp Thr Val Thr Leu 115 120 125
- Glu Val Leu Val Ala Pro Ala Val Pro Ser Cys Glu Val Pro Ser Ser 130 135 140
- Ala Leu Ser Gly Thr Val Val Glu Leu Arg Cys Gln Asp Lys Glu Gly 145 150 155 160
- Asn Pro Ala Pro Glu Tyr Thr Trp Phe Lys Asp Gly Ile Arg Leu Leu 165 170 175
- Glu Asn Pro Arg Leu Gly Ser Gln Ser Thr Asn Ser Ser Tyr Thr Met 180 185 190
- Asn Thr Lys Thr Gly Thr Leu Gln Phe Asn Thr Val Ser Lys Leu Asp 195 200 205
- Thr Gly Glu Tyr Ser Cys Glu Ala Arg Asn Ser Val Gly Tyr Arg Arg 210 215 220
- Cys Pro Gly Lys Arg Met Gln Val Asp Asp Leu Asn Ile Ser Gly Ile 225 230 235 240
- Ile Ala Ala Val Val Val Ala Leu Val Ile Ser Val Cys Gly Leu 245 250 255
- Gly Val Cys Tyr Ala Gln Arg Lys Gly Tyr Phe Ser Lys Glu Thr Ser 260 265 270
- Phe Gln Lys Ser Asn Ser Ser Ser Lys Ala Thr Thr Met Ser Glu Asn 275 280 285
- Val Gln Trp Leu Thr Pro Val Ile Pro Ala Leu Trp Lys Ala Ala Ala 290 295 300
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Ser Val Phe His Ser Val Gly Asn Leu Glu Thr Leu Ile Leu Asp Ser 355 360 365

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Trp Arg Leu Asn Phe Asn Arg Gln Gln Pro Thr Cys Ala Thr Pro Glu 385 390 395 400

Phe Val Gln Gly Lys Glu Phe Lys Asp Phe Pro Asp Val Leu Leu Pro 405 410 415

Asn Tyr Phe Thr Cys Arg Arg Ala Arg Ile Arg Asp Arg Lys Ala Gln
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Asp Gly Asp Pro Pro Pro Ala Ile Leu Trp Leu Ser Pro Arg Lys His 450 455 460

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Thr Leu Glu Val Arg Tyr Ala Gln Val Gln Asp Asn Gly Thr Tyr Leu 485 490 495

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- Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu Gln Trp Ser Asn
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- Asn Arg Ile Gln Leu Val Thr Ser Thr Pro His Glu Leu Ser Ile Ser 85 90 95
- Ile Ser Asn Val Ala Leu Ala Asp Glu Gly Glu Tyr Thr Cys Ser Ile 100 105 110
- Phe Thr Met Pro Val Arg Thr Ala Lys Ser Leu Val Thr Val Leu Gly
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- Ile Pro Gln Lys Pro Ile Ile Thr Gly Tyr Lys Ser Ser Leu Arg Glu 130 135 140
- Lys Asp Thr Ala Thr Leu Asn Cys Gln Ser Ser Gly Ser Lys Pro Ala 145 150 155 160
- Ala Arg Leu Thr Trp Arg Lys Gly Asp Gln Glu Leu His Gly Glu Pro 165 170 175
- Thr Arg Ile Gln Glu Asp Pro Asn Gly Lys Thr Phe Thr Val Ser Ser 180 185 190
- Ser Val Thr Phe Gln Val Thr Arg Glu Asp Asp Gly Ala Ser Ile Val 195 200 205
- Cys Ser Val Asn His Glu Ser Leu Lys Gly Ala Asp Arg Ser Thr Ser 210 215 220
- Gln Arg Ile Glu Val Leu Tyr Thr Pro Thr Ala Met Ile Arg Pro Asp 225 230 235 240
- Pro Pro His Pro Arg Glu Gly Gln Lys Leu Leu His Cys Glu Gly 245 250 255
- Arg Gly Asn Pro Val Pro Gln Gln Tyr Leu Trp Glu Lys Glu Gly Ser 260 265 270
- Val Pro Pro Leu Lys Met Thr Gln Glu Ser Ala Leu Ile Phe Pro Phe 275 280 285

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                                     330
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His Tyr Leu Ile Arg His Lys Gly Thr Tyr Leu Thr His Glu Ala Lys
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## oligonucleotide probe

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| agagaagate tgtteetgea atgagataga aggggaeeta caegtagaet gtgaaaaaaa  |      |
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| ccgagtggat tctagtctcc cggcgccccc tgcccaagaa gagacctttg ctcctggacc  | 1020 |
| cctgccaact cctttcaaga caaatgggca agaggatcat gccacaccag ggtctgctcc  | 1080 |

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Leu His Gly Asn Ser Leu Thr Arg Leu Phe Pro Asn Glu Phe Ala Asn
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Phe Tyr Asn Ala Val Ser Leu His Met Glu Asn Asn Gly Leu His Glu

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- Ile Asn Asn Asn Lys Ile Lys Ser Phe Arg Lys Gln Thr Phe Leu Gly
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- Ile Asp Pro Gly Ala Phe Gln Asp Leu Asn Lys Leu Glu Val Leu Ile 145 150 155 160
- Leu Asn Asp Asn Leu Ile Ser Thr Leu Pro Ala Asn Val Phe Gln Tyr 165 170 175
- Val Pro Ile Thr His Leu Asp Leu Arg Gly Asn Arg Leu Lys Thr Leu 180 185 190
- Pro Tyr Glu Glu Val Leu Glu Gln Ile Pro Gly Ile Ala Glu Ile Leu 195 200 205
- Leu Glu Asp Asn Pro Trp Asp Cys Thr Cys Asp Leu Leu Ser Leu Lys 210 215 220
- Glu Trp Leu Glu Asn Ile Pro Lys Asn Ala Leu Ile Gly Arg Val Val 225 230 235 240
- Cys Glu Ala Pro Thr Arg Leu Gln Gly Lys Asp Leu Asn Glu Thr Thr 245 250 255
- Glu Gln Asp Leu Cys Pro Leu Lys Asn Arg Val Asp Ser Ser Leu Pro 260 265 270
- Ala Pro Pro Ala Gln Glu Glu Thr Phe Ala Pro Gly Pro Leu Pro Thr 275 280 285
- Pro Phe Lys Thr Asn Gly Gln Glu Asp His Ala Thr Pro Gly Ser Ala 290 295 300
- Pro Asn Gly Gly Thr Lys Ile Pro Gly Asn Trp Gln Ile Lys Ile Arg 305 310 315 320
- Pro Thr Ala Ala Ile Ala Thr Gly Ser Ser Arg Asn Lys Pro Leu Ala 325 330 335
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- Ser Gly Leu Lys Met Asn Cys Asn Asn Arg Asn Val Ser Ser Leu Ala 355 360 365
- Asp Leu Lys Pro Lys Leu Ser Asn Val Gln Glu Leu Phe Leu Arg Asp 370 375 380

- Asn Lys Ile His Ser Ile Arg Lys Ser His Phe Val Asp Tyr Lys Asn 385 390 395 400
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- Pro Val Asn Phe Phe Arg Lys Asp Phe Met Leu Leu Ser Asn Asp Glu 565 570 575
- Ile Cys Pro Gln Leu Tyr Ala Arg Ile Ser Pro Thr Leu Thr Ser His 580 585 590
- Ser Lys Asn Ser Thr Gly Leu Ala Glu Thr Gly Thr His Ser Asn Ser 595 600 605
- Tyr Leu Asp Thr Ser Arg Val Ser Ile Ser Val Leu Val Pro Gly Leu 610 615 620
- Leu Leu Val Phe Val Thr Ser Ala Phe Thr Val Val Gly Met Leu Val 625 630 635 640
- Phe Ile Leu Arg Asn Arg Lys Arg Ser Lys Arg Arg Asp Ala Asn Ser 645 650 655
- Ser Ala Ser Glu Ile Asn Ser Leu Gln Thr Val Cys Asp Ser Ser Tyr

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- Asp Asn Ser Val Thr Ser Ile Pro Glu Ile Pro Arg Trp Gly Ser Gln

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| Thr          | Ile<br>370                       | Thr           | Pro        | Ser        | Gly        | Ser<br>375 | Val        | Ile        | Ser        | Lys        | Phe<br>380 | Asn        | Ser        | Thr        | Thr        |    |
| Ser<br>385   | Ser                              | Ala           | Thr        | Pro        | Gln<br>390 | Ala        | Phe        | Asp        | Ser        | Ser<br>395 | Ser        | Ala        | Val        | Val        | Phe<br>400 |    |
| Ile          | Phe                              | Val           | Ser        | Thr<br>405 | Ala        | Val        | Val        | Val        | Leu<br>410 | Val        | Ile        | Leu        | Thr        | Met<br>415 | Thr        |    |
| Val          | Leu                              | Gly           | Leu<br>420 | Val        | Lys        | Leu        | Cys        | Phe<br>425 | His        | Glu        | Ser        | Pro        | Ser<br>430 | Ser        | Gln        |    |
| Pro          | Arg                              | Lys<br>435    | Glu        | Ser        | Met        | Gly        | Pro<br>440 | Pro        | Gly        | Leu        | Glu        | Ser<br>445 | Asp        | Pro        | Glu        |    |
| Pro          | Ala<br>450                       | Ala           | Leu        | Gly        | Ser        | Ser<br>455 | Ser        | Ala        | His        | Cys        | Thr<br>460 | Asn        | Asn        | Gly        | Val        |    |
| Lys<br>465   | Val                              | Gly           | Asp        | Cys        | Asp<br>470 | Leu        | Arg        | Asp        | Arg        | Ala<br>475 | Glu        | Gly        | Ala        | Leu        | Leu<br>480 |    |
| Ala          | Glu                              | Ser           | Pro        | Leu<br>485 | Gly        | Ser        | Ser        | Asp        | Ala<br>490 |            |            |            |            |            |            |    |
| <211<br><212 | 0> 9'<br>L> 24<br>2> DI<br>3> A: | 4             | icia       | l Se       | quen       | ce         |            |            |            |            |            |            |            |            |            |    |
| <220         | )>                               |               |            |            |            |            |            |            |            |            |            |            |            |            |            |    |
|              | 3 > De                           | escr:<br>ligo |            |            |            |            | cial       | Sequ       | uenc       | e: S       | ynth       | etic       |            |            |            |    |
|              | )> 9'<br>aagg                    | 7<br>aga 1    | tgcg       | atgc       | ca c       | ctg        |            |            |            |            |            |            |            |            |            | 24 |
| <212<br><212 | 0> 9:<br>L> 2:<br>2> D:<br>3> A: | 0             | icia       | l Se       | quen       | ce         |            |            |            |            |            |            |            |            |            |    |
| <220<br><220 | 3 > D                            | escr<br>ligo  |            |            |            |            | cial       | Seq        | uenc       | e: S       | ynth       | etic       |            |            |            |    |
|              | )> 9<br>ccag                     | 8<br>tgg (    | ggaa       | ggac       | ag         |            |            |            |            |            |            |            |            |            |            | 20 |

| <210> 99 <211> 20 <212> DNA <213> Artificial Sequence                           |    |
|---|----|
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 99 acagagcaga gggtgccttg  | 20 |
| <210> 100<br><211> 24<br><212> DNA<br><213> Artificial Sequence                 |    |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 100<br>tcagggacaa gtggtgtctc tccc   | 24 |
| <210> 101<br><211> 24<br><212> DNA<br><213> Artificial Sequence                 |    |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 101<br>tcagggaagg agtgtgcagt tctg   | 24 |
| <210> 102<br><211> 50<br><212> DNA<br><213> Artificial Sequence                 |    |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 102<br>acageteceg ateteagtta ettgeatege ggacgaaate ggegeteget             | 50 |
| <210> 103<br><211> 2026<br><212> DNA<br><213> Homo sapiens                      |    |

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tatcccccgg ctacctgggc cgccccgcgg cggtgcgcgc gtgagaggga gcgcgcgggc 180
agccgagcgc cggtgtgagc cagcgctgct gccagtgtga gcggcggtgt gagcgcggtg 240
ggtgcggagg ggcgtgtgtg ccggcgcgcg cgccgtgggg tgcaaacccc gagcgtctac 300
gctgccatga ggggcgcgaa cgcctgggcg ccactctgcc tgctgctggc tgccgccacc 360
cagetetege ggeageagte eccagagaga cetgttttea catgtggtgg cattettaet 420
ggagagtctg gatttattgg cagtgaaggt tttcctggag tgtaccctcc aaatagcaaa 480
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gtcacttgtg tgtggcacat tgtagcccca aagaatcagc ttatagaatt aaagtttgag 900
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attttagaat tgagttgtgt gaagatgtca aaaaaagatt ttagaagtgc aatatttata 1920
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<211> 415
<212> PRT
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             20
Cys Gly Gly Ile Leu Thr Gly Glu Ser Gly Phe Ile Gly Ser Glu Gly
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Phe Pro Gly Val Tyr Pro Pro Asn Ser Lys Cys Thr Trp Lys Ile Thr

- Val Pro Glu Gly Lys Val Val Leu Asn Phe Arg Phe Ile Asp Leu 65 70 75 80
- Glu Ser Asp Asn Leu Cys Arg Tyr Asp Phe Val Asp Val Tyr Asn Gly
  85 90 95
- His Ala Asn Gly Gln Arg Ile Gly Arg Phe Cys Gly Thr Phe Arg Pro
- Gly Ala Leu Val Ser Ser Gly Asn Lys Met Met Val Gln Met Ile Ser
- Asp Ala Asn Thr Ala Gly Asn Gly Phe Met Ala Met Phe Ser Ala Ala 130 135 140
- Glu Pro Asn Glu Arg Gly Asp Gln Tyr Cys Gly Gly Leu Leu Asp Arg 145 150 155 160
- Pro Ser Gly Ser Phe Lys Thr Pro Asn Trp Pro Asp Arg Asp Tyr Pro 165 170 175
- Ala Gly Val Thr Cys Val Trp His Ile Val Ala Pro Lys Asn Gln Leu 180 185 190
- Ile Glu Leu Lys Phe Glu Lys Phe Asp Val Glu Arg Asp Asn Tyr Cys
- Arg Tyr Asp Tyr Val Ala Val Phe Asn Gly Glu Val Asn Asp Ala 210 215 220
- Arg Arg Ile Gly Lys Tyr Cys Gly Asp Ser Pro Pro Ala Pro Ile Val 225 230 235 240
- Ser Glu Arg Asn Glu Leu Leu Ile Gln Phe Leu Ser Asp Leu Ser Leu 245 250 255
- Thr Ala Asp Gly Phe Ile Gly His Tyr Ile Phe Arg Pro Lys Lys Leu 260 265 270
- Pro Thr Thr Glu Gln Pro Val Thr Thr Thr Phe Pro Val Thr Thr 275 280 285
- Gly Leu Lys Pro Thr Val Ala Leu Cys Gln Gln Lys Cys Arg Arg Thr 290 295 300
- Gly Thr Leu Glu Gly Asn Tyr Cys Ser Ser Asp Phe Val Leu Ala Gly 305 310 315 320
- Thr Val Ile Thr Thr Ile Thr Arg Asp Gly Ser Leu His Ala Thr Val 325 330 335
- Ser Ile Ile Asn Ile Tyr Lys Glu Gly Asn Leu Ala Ile Gln Gln Ala

|                               |                    | 340  |            |            |            |            | 345  |            |            |            |            | 350 |            |            |    |
|-------------------------------|--------------------|------|------------|------------|------------|------------|------|------------|------------|------------|------------|-----|------------|------------|----|
| Gly Ly                        | s Asn<br>355       |      | Ser        | Ala        | Arg        | Leu<br>360 | Thr  | Val        | Val        | Cys        | Lys<br>365 | Gln | Cys        | Pro        |    |
| Leu Le                        |                    | Arg  | Gly        | Leu        | Asn<br>375 | Tyr        | Ile  | Ile        | Met        | Gly<br>380 | Gln        | Val | Gly        | Glu        |    |
| Asp Gl<br>385                 | y Arg              | Gly  | Lys        | Ile<br>390 | Met        | Pro        | Asn  | Ser        | Phe<br>395 | Ile        | Met        | Met | Phe        | Lys<br>400 |    |
| Thr Ly                        | s Asn              | Gln  | Lys<br>405 | Leu        | Leu        | Asp        | Ala  | Leu<br>410 | Lys        | Asn        | Lys        | Gln | Cys<br>415 |            |    |
| <210> <211> <212> <213> <223> | 22<br>DNA<br>Artif | ipti | on o       | f Ar       | tifi       | cial       | Seq  | uence      | e: S       | ynth       | etic       |     |            |            |    |
| <400><br>ccgatt               | 105                |      |            |            |            |            |      |            |            |            |            |     |            |            | 22 |
| <210><211><211><212><213>     | 22<br>DNA          | icia | l Se       | quen       | ce         |            |      |            |            |            |            |     |            |            |    |
| <220><br><223>                | Descr<br>oligo     |      |            |            |            |            | Seq  | uenc       | e: S       | ynth       | etic       |     |            |            |    |
| <400><br>gtcaag               |                    | cctc | caca       | at a       | С          |            |      |            |            |            |            |     |            |            | 22 |
| <210><211><212><212><213>     | 45<br>DNA          | icia | l Se       | quen       | ce         |            |      |            |            |            |            |     |            |            |    |
| <220><br><223>                | Desci<br>oligo     |      |            |            |            |            | Seq  | uenc       | e: S       | ynth       | etic       | !   |            |            |    |
| <400><br>gtgtac               |                    | gcca | ıtgcc      | aa t       | ggcc       | agcg       | c at | tggc       | cgct       | tct        | gt         |     |            |            | 45 |
| <210><211><211>               | 1838               |      |            |            |            |            |      |            |            |            |            |     |            |            |    |

## <213> Homo sapiens

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aagggcctag tcccagctgt gctctggggc ctcagcctct tcctcaacct cccaggacct 180
atotggotoc agocototoc acotococag tottotococ egeotoagoc coatoegtgt 240
catacctgcc ggggactggt tgacagcttt aacaagggcc tggagagaac catccgggac 300
aactttggag gtggaaacac tgcctgggag gaagagaatt tgtccaaata caaagacagt 360
gagacccgcc tggtagaggt gctggagggt gtgtgcagca agtcagactt cgagtgccac 420
cqcctqctqq aqctqagtga ggagctggtg gagagctggt ggtttcacaa gcagcaggag 480
gccccggacc tcttccagtg gctgtgctca gattccctga agctctgctg ccccgcaggc 540
accttcgggc cctcctgcct tccctgtcct gggggaacag agaggccctg cggtggctac 600
gggcagtgtg aaggagaagg gacacgaggg ggcagcgggc actgtgactg ccaagccggc 660
tacqqqqqtq aggcctgtgg ccagtgtggc cttggctact ttgaggcaga acgcaacgcc 720
agccatctgg tatgttcggc ttgttttggc ccctgtgccc gatgctcagg acctgaggaa 780
tcaaactgtt tgcaatgcaa gaagggctgg gccctgcatc acctcaagtg tgtagacatt 840
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ggtcgctgta agaagtgtag ccctggctat cagcaggtgg gctccaagtg tctcgatgtg 1020
gatgagtgtg agacagaggt gtgtccggga gagaacaagc agtgtgaaaa caccgagggc 1080
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cagcagatgt tetttggcat catcatetgt gcaetggcca egetggetge taagggegae 1260
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<210> 109
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<213> Homo sapiens

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Pro Pro Gln Ser Ser Pro Pro Gln Pro His Pro Cys His Thr

Cys Arg Gly Leu Val Asp Ser Phe Asn Lys Gly Leu Glu Arg Thr Ile
50 55 60

<sup>&</sup>lt;211> 420

<sup>&</sup>lt;212> PRT

- Arg Asp Asn Phe Gly Gly Gly Asn Thr Ala Trp Glu Glu Glu Asn Leu 65 70 75 80
- Ser Lys Tyr Lys Asp Ser Glu Thr Arg Leu Val Glu Val Leu Glu Gly
  85 90 95
- Val Cys Ser Lys Ser Asp Phe Glu Cys His Arg Leu Leu Glu Leu Ser 100 105 110
- Glu Glu Leu Val Glu Ser Trp Trp Phe His Lys Gln Gln Glu Ala Pro 115 120 125
- Asp Leu Phe Gln Trp Leu Cys Ser Asp Ser Leu Lys Leu Cys Cys Pro 130 140
- Ala Gly Thr Phe Gly Pro Ser Cys Leu Pro Cys Pro Gly Gly Thr Glu
  145 150 155 160
- Arg Pro Cys Gly Gly Tyr Gly Gln Cys Glu Gly Glu Gly Thr Arg Gly
  165 170 175
- Gly Ser Gly His Cys Asp Cys Gln Ala Gly Tyr Gly Glu Ala Cys 180 185 190
- Gly Gln Cys Gly Leu Gly Tyr Phe Glu Ala Glu Arg Asn Ala Ser His 195 200 205
- Leu Val Cys Ser Ala Cys Phe Gly Pro Cys Ala Arg Cys Ser Gly Pro 210 215 220
- Glu Glu Ser Asn Cys Leu Gln Cys Lys Lys Gly Trp Ala Leu His His 225 230 235 240
- Leu Lys Cys Val Asp Ile Asp Glu Cys Gly Thr Glu Gly Ala Asn Cys 245 250 255
- Gly Ala Asp Gln Phe Cys Val Asn Thr Glu Gly Ser Tyr Glu Cys Arg 260 265 270
- Asp Cys Ala Lys Ala Cys Leu Gly Cys Met Gly Ala Gly Pro Gly Arg 275 280 285
- Cys Lys Lys Cys Ser Pro Gly Tyr Gln Gln Val Gly Ser Lys Cys Leu 290 295 300
- Asp Val Asp Glu Cys Glu Thr Glu Val Cys Pro Gly Glu Asn Lys Gln 305 310 315 320
- Cys Glu Asn Thr Glu Gly Gly Tyr Arg Cys Ile Cys Ala Glu Gly Tyr 325 330 335
- Lys Gln Met Glu Gly Ile Cys Val Lys Glu Gln Ile Pro Glu Ser Ala 340 345 350

| Gly Phe Phe Ser Glu Met Thr Glu Asp Glu Leu Val Val Leu G<br>355 360 365        | ln Gln         |
|---|----------------|
| Met Phe Phe Gly Ile Ile Ile Cys Ala Leu Ala Thr Leu Ala A<br>370 375 380        | la Lys         |
| Gly Asp Leu Val Phe Thr Ala Ile Phe Ile Gly Ala Val Ala A<br>385 390 395        | ala Met<br>400 |
| Thr Gly Tyr Trp Leu Ser Glu Arg Ser Asp Arg Val Leu Glu G<br>405 410 4          | ly Phe<br>15   |
| Ile Lys Gly Arg 420   |                |
| <210> 110<br><211> 50<br><212> DNA<br><213> Artificial Sequence                 |                |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |                |
| <400> 110<br>cctggctatc agcaggtggg ctccaagtgt ctcgatgtgg atgagtgtga             | 50             |
| <210> 111<br><211> 22<br><212> DNA<br><213> Artificial Sequence                 |                |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |                |
| <400> 111<br>attctgcgtg aacactgagg gc   | 22             |
| <210> 112<br><211> 22<br><212> DNA<br><213> Artificial Sequence                 |                |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |                |
| <400> 112<br>atctgcttgt agccctcggc ac   | 22             |
| <210> 113   |                |

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<212> DNA
<213> Homo sapiens
<220>
<221> modified_base
<222> (1461)
<223> a, t, c or g
<400> 113
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cggggccgcc ctgaccgggg agcagctcct gggcagcctg ctgcggcagc tgcagctcaa 180
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cagccagage ttccgagagg tggccggcag gttcctggcg ttggaggcca gcacacacct 360
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geggetette caggageegg teeccaagge egegetgeae aggeaeggge ggetgteece 480
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gctacaggtg teggtgcaga gggagcatet gggceegetg gegteeggeg eccaeaaget 720
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caccetggae ettggggaet atggagetea gggegaetgt gaeeetgaag eaccaatgae 840
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aaagtcctcc accaccactc tggacctaag acctggggtt aagtgtgggt tgtgcatccc 1560
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<210> 114
<211> 366
<212> PRT
<213> Homo sapiens
<400> 114
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Ser Pro Gly Ala Ala Leu Thr Gly Glu Gln Leu Leu Gly Ser Leu Leu
             20
                                  25
Arg Gln Leu Gln Leu Lys Glu Val Pro Thr Leu Asp Arg Ala Asp Met
                              40
         35
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- Glu Glu Leu Val Ile Pro Thr His Val Arg Ala Gln Tyr Val Ala Leu 50 55 60
- Leu Gln Arg Ser His Gly Asp Arg Ser Arg Gly Lys Arg Phe Ser Gln 65 70 75 80
- Ser Phe Arg Glu Val Ala Gly Arg Phe Leu Ala Leu Glu Ala Ser Thr 85 90 95
- His Leu Leu Val Phe Gly Met Glu Gln Arg Leu Pro Pro Asn Ser Glu 100 105 110
- Leu Val Gln Ala Val Leu Arg Leu Phe Gln Glu Pro Val Pro Lys Ala 115 120 125
- Ala Leu His Arg His Gly Arg Leu Ser Pro Arg Ser Ala Arg Ala Arg 130 135 140
- Val Thr Val Glu Trp Leu Arg Val Arg Asp Asp Gly Ser Asn Arg Thr 145 150 155 160
- Ser Leu Ile Asp Ser Arg Leu Val Ser Val His Glu Ser Gly Trp Lys 165 170 175
- Ala Phe Asp Val Thr Glu Ala Val Asn Phe Trp Gln Gln Leu Ser Arg 180 185 190
- Pro Arg Gln Pro Leu Leu Gln Val Ser Val Gln Arg Glu His Leu 195 200 205
- Gly Pro Leu Ala Ser Gly Ala His Lys Leu Val Arg Phe Ala Ser Gln 210 215 220
- Gly Ala Pro Ala Gly Leu Gly Glu Pro Gln Leu Glu Leu His Thr Leu 225 230 235 240
- Asp Leu Gly Asp Tyr Gly Ala Gln Gly Asp Cys Asp Pro Glu Ala Pro 245 250 255
- Met Thr Glu Gly Thr Arg Cys Cys Arg Gln Glu Met Tyr Ile Asp Leu 260 265 270
- Gln Gly Met Lys Trp Ala Glu Asn Trp Val Leu Glu Pro Pro Gly Phe 275 280 285
- Leu Ala Tyr Glu Cys Val Gly Thr Cys Arg Gln Pro Pro Glu Ala Leu 290 295 300
- Ala Phe Lys Trp Pro Phe Leu Gly Pro Arg Gln Cys Ile Ala Ser Glu 305 310 315 320
- Thr Asp Ser Leu Pro Met Ile Val Ser Ile Lys Glu Gly Gly Arg Thr 325 330 335

| Arg Pro Gln Val Val Ser Leu Pro Asn Met Arg Val Gln Lys Cys Ser 340 345 350   |     |
|---|-----|
| Cys Ala Ser Asp Gly Ala Leu Val Pro Arg Arg Leu Gln Pro<br>355 360 365  |     |
| <210> 115<br><211> 21<br><212> DNA<br><213> Artificial Sequence   |     |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe   |     |
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| <210> 116<br><211> 22<br><212> DNA<br><213> Artificial Sequence   |     |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe   |     |
| <400> 116<br>ataggagttg aagcagcgct gc   | 22  |
| <210> 117<br><211> 45<br><212> DNA<br><213> Artificial Sequence   |     |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe   |     |
| <400> 117<br>tgtgtggaca tagacgagtg ccgctaccgc tactgccagc accgc  | 45  |
| <210> 118<br><211> 1857<br><212> DNA<br><213> Homo sapiens  |     |
| <400> 118 gtctgttccc aggagtcctt cggcggctgt tgtgtcagtg gcctgatcgc gatggggaca aaggcgcaag tcgagaggaa actgttgtgc ctcttcatat tggcgatcct gttgtgctcc ctggcattgg gcagtgttac agtgcactct tctgaacctg aagtcagaat tcctgagaat | 120 |

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gtcaagetca tegtgettgt geetceatee aageetacag ttaacateee eteetetgee 480
accattggga accgggcagt gctgacatgc tcagaacaag atggttcccc accttctgaa 540
tacacctggt tcaaagatgg gatagtgatg cctacgaatc ccaaaagcac ccgtgccttc 600
agcaactett cetatgteet gaateecaca acaggagage tggtetttga teecetgtea 660
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Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu
                                                  45
         35
Ser Cys Ala Tyr Ser Gly Phe Ser Ser Pro Arg Val Glu Trp Lys Phe
                          55
Asp Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr
                                          75
 65
                      70
Ala Ser Tyr Glu Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr Phe
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90

Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser

Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val 115 120 125

Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala Thr 130 135 140

Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro 145 150 155 160

Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn 165 170 175

Pro Lys Ser Thr Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro 180 185 190

Thr Thr Gly Glu Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly 195 200 205

Glu Tyr Ser Cys Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser 210 215 220

Asn Ala Val Arg Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val 225 230 235 240

Ala Ala Val Leu Val Thr Leu Ile Leu Leu Gly Ile Leu Val Phe Gly 245 250 255

Ile Trp Phe Ala Tyr Ser Arg Gly His Phe Asp Arg Thr Lys Lys Gly

Thr Ser Ser Lys Lys Val Ile Tyr Ser Gln Pro Ser Ala Arg Ser Glu 275 280 285

Gly Glu Phe Lys Gln Thr Ser Ser Phe Leu Val 290 295

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<212> DNA

<213> Artificial Sequence

<220>

<400> 120

tcgcggagct gtgttctgtt tccc

<210> 121

<211> 50

| <212> DNA<br><213> Artificial Sequence  |    |
|---|----|
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 121<br>tgatcgcgat ggggacaaag gcgcaagctc gagaggaaac tgttgtgcct             | 50 |
| <210> 122<br><211> 20<br><212> DNA<br><213> Artificial Sequence                 |    |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 122<br>acacctggtt caaagatggg  | 20 |
| <210> 123<br><211> 24<br><212> DNA<br><213> Artificial Sequence                 |    |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 123<br>taggaagagt tgctgaaggc acgg   | 24 |
| <210> 124<br><211> 20<br><212> DNA<br><213> Artificial Sequence                 |    |
| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
| <400> 124<br>ttgccttact caggtgctac  | 20 |
| <210> 125<br><211> 20<br><212> DNA<br><213> Artificial Sequence                 |    |
| <220> <223> Description of Artificial Sequence: Synthetic                       |    |

## oligonucleotide probe

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<400> 125
                                                                  2.0
actcagcagt ggtaggaaag
<210> 126
<211> 1210
<212> DNA
<213> Homo sapiens
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qcctqqaqqc cqccqcqaqc ccgctttcca ccccgacctc tgcccaggcc gcaggcccca 180
gctcaggctc gtgcccaccc accaagttcc agtgccgcac cagtggctta tgcgtgcccc 240
tcacctggcg ctgcgacagg gacttggact gcagcgatgg cagcgatgag gaggagtgca 300
ggattgagcc atgtacccag aaagggcaat gcccaccgcc ccctggcctc ccctgcccct 360
qcaccqqcqt cagtgactgc tctgggggaa ctgacaagaa actgcgcaac tgcagccgcc 420
tggcctgcct agcaggcgag ctccgttgca cgctgagcga tgactgcatt ccactcacgt 480
qqcqctqcga cggccaccca gactgtcccg actccagcga cgagctcggc tgtggaacca 540
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                                                      30
Ala Ala Ala Ser Pro Leu Ser Thr Pro Thr Ser Ala Gln Ala Ala Gly
         35
                             40
                                                 45
Pro Ser Ser Gly Ser Cys Pro Pro Thr Lys Phe Gln Cys Arg Thr Ser
                         55
Gly Leu Cys Val Pro Leu Thr Trp Arg Cys Asp Arg Asp Leu Asp Cys
                                         75
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Ser Asp Gly Ser Asp Glu Glu Glu Cys Arg Ile Glu Pro Cys Thr Gln
85 90 95

Lys Gly Gln Cys Pro Pro Pro Pro Gly Leu Pro Cys Pro Cys Thr Gly
100 105 110

Val Ser Asp Cys Ser Gly Gly Thr Asp Lys Lys Leu Arg Asn Cys Ser 115 120 125

Arg Leu Ala Cys Leu Ala Gly Glu Leu Arg Cys Thr Leu Ser Asp Asp 130 135 140

Cys Ile Pro Leu Thr Trp Arg Cys Asp Gly His Pro Asp Cys Pro Asp 145 150 155 160

Ser Ser Asp Glu Leu Gly Cys Gly Thr Asn Glu Ile Leu Pro Glu Gly 165 170 175

Asp Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val Thr Ser 180 185 190

Leu Arg Asn Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val
195 200 205

Pro Ser Val Gly Asn Ala Thr Ser Ser Ser Ala Gly Asp Gln Ser Gly 210 215 220

Ser Pro Thr Ala Tyr Gly Val Ile Ala Ala Ala Ala Val Leu Ser Ala 225 230 235 240

Ser Leu Val Thr Ala Thr Leu Leu Leu Leu Ser Trp Leu Arg Ala Gln 245 250 255

Glu Arg Leu Arg Pro Leu Gly Leu Leu Val Ala Met Lys Glu Ser Leu 260 265 270

Leu Leu Ser Glu Gln Lys Thr Ser Leu Pro 275 280

<210> 128

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<400> 128

aagttccagt gccgcaccag tggc

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<211> 50
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<210> 131
<211> 1843
<212> DNA
<213> Homo sapiens
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<221> modified base
<222> (1837)
<223> a, t, c or g
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cagactettg caagetggat geeetetgtg gatgaaagat gtateatgga atgaaceega 180
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gctggatccc aagtgataat tccatctgtg tgcaagaaga ttgccgtatc cctcaaatcg 540
aagatgctga gattcataac aagacatata gacatggaga gaagctaatc atcacttgtc 600
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atggctatgt aaacatetet gageteeaga eeteetteee ggtggggaet gtgateteet 780
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agcaaacgtg gcccagcacc catgagaccc tcctgaccac gtggaagatt gtggcgttca 1140
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Ile Leu Trp Phe Gln Leu Ala Leu Cys Phe Gly Pro Ala Gln Leu Thr
                             40
Gly Gly Phe Asp Asp Leu Gln Val Cys Ala Asp Pro Gly Ile Pro Glu
Asn Gly Phe Arg Thr Pro Ser Gly Gly Val Phe Phe Glu Gly Ser Val
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Ala Arg Phe His Cys Gln Asp Gly Phe Lys Leu Lys Gly Ala Thr Lys
                                     90
                 85
Arg Leu Cys Leu Lys His Phe Asn Gly Thr Leu Gly Trp Ile Pro Ser
                                                     110
                                105
Asp Asn Ser Ile Cys Val Gln Glu Asp Cys Arg Ile Pro Gln Ile Glu
                            120
                                                125
Asp Ala Glu Ile His Asn Lys Thr Tyr Arg His Gly Glu Lys Leu Ile
    130
                        135
Ile Thr Cys His Glu Gly Phe Lys Ile Arg Tyr Pro Asp Leu His Asn
                                        155
                    150
Met Val Ser Leu Cys Arg Asp Asp Gly Thr Trp Asn Asn Leu Pro Ile
                                                         175
                165
Cys Gln Gly Cys Leu Arg Pro Leu Ala Ser Ser Asn Gly Tyr Val Asn
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|            |            |            | 180        |            |            |            |            | 185        |            |            |            |            | 190        |            |            |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ile        | Ser        | Glu<br>195 | Leu        | Gln        | Thr        | Ser        | Phe<br>200 | Pro        | Val        | Gly        | Thr        | Val<br>205 | Ile        | Ser        | Tyr        |
| Arg        | Cys<br>210 | Phe        | Pro        | Gly        | Phe        | Lys<br>215 | Leu        | Asp        | Gly        | Ser        | Ala<br>220 | Tyr        | Leu        | Glu        | Cys        |
| Leu<br>225 | Gln        | Asn        | Leu        | Ile        | Trp<br>230 | Ser        | Ser        | Ser        | Pro        | Pro<br>235 | Arg        | Cys        | Leu        | Ala        | Leu<br>240 |
| Glu        | Ala        | Gln        | Val        | Cys<br>245 | Pro        | Leu        | Pro        | Pro        | Met<br>250 | Val        | Ser        | His        | Gly        | Asp<br>255 | Phe        |
| Val        | Cys        | His        | Pro<br>260 | Arg        | Pro        | Cys        | Glu        | Arg<br>265 | Tyr        | Asn        | His        | Gly        | Thr<br>270 | Val        | Val        |
| Glu        | Phe        | Tyr<br>275 | Cys        | Asp        | Pro        | Gly        | Tyr<br>280 | Ser        | Leu        | Thr        | Ser        | Asp<br>285 | Tyr        | Lys        | Tyr        |
| Ile        | Thr<br>290 | Cys        | Gln        | Tyr        | Gly        | Glu<br>295 | Trp        | Phe        | Pro        | Ser        | Tyr<br>300 | Gln        | Val        | Tyr        | Cys        |
| Ile<br>305 | Lys        | Ser        | Glu        | Gln        | Thr<br>310 | Trp        | Pro        | Ser        | Thr        | His<br>315 | Glu        | Thr        | Leu        | Leu        | Thr<br>320 |
| Thr        | Trp        | Lys        | Ile        | Val<br>325 | Ala        | Phe        | Thr        | Ala        | Thr<br>330 | Ser        | Val        | Leu        | Leu        | Val<br>335 | Leu        |
| Leu        | Leu        | Val        | Ile<br>340 | Leu        | Ala        | Arg        | Met        | Phe<br>345 | Gln        | Thr        | Lys        | Phe        | Lys<br>350 | Ala        | His        |
| Phe        | Pro        | Pro<br>355 | Arg        | Gly        | Pro        | Pro        | Arg<br>360 | Ser        | Ser        | Ser        | Ser        | Asp<br>365 | Pro        | Asp        | Phe        |
| Val        | Val<br>370 | Val        | Asp        | Gly        | Val        | Pro<br>375 | Val        | Met        | Leu        | Pro        | Ser<br>380 | Tyr        | Asp        | Glu        | Ala        |
| Val<br>385 | Ser        | Gly        | Gly        | Leu        | Ser<br>390 | Ala        | Leu        | Gly        | Pro        | Gly<br>395 | Tyr        | Met        | Ala        | Ser        | Val<br>400 |
| Gly        | Gln        | Gly        | Cys        | Pro<br>405 | Leu        | Pro        | Val        | Asp        | Asp<br>410 | Gln        | Ser        | Pro        | Pro        | Ala<br>415 | Tyr        |
| Pro        | Gly        | Ser        | Gly<br>420 | Asp        | Thr        | Asp        | Thr        | Gly<br>425 | Pro        | Gly        | Glu        | Ser        | Glu<br>430 | Thr        | Cys        |
| Asp        | Ser        | Val<br>435 | Ser        | Gly        | Ser        | Ser        | Glu<br>440 | Leu        | Leu        | Gln        | Ser        | Leu<br>445 | Tyr        | Ser        | Pro        |
| Pro        | Arg        | _          | Gln        | Glu        | Ser        | Thr<br>455 | His        | Pro        | Ala        | Ser        | Asp<br>460 |            | Pro        | Asp        | Ile        |

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                                         475
465
His Ala His Trp Val Leu Phe Leu Arg Asn
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<400> 133
                                                                    23
atctcctatc gctgctttcc cgg
<210> 134
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<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 134
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<212> DNA
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<223> Description of Artificial Sequence: Synthetic
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<210> 136
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<212> DNA
<213> Homo sapiens
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ccgtagcgcc cgagtgtcgg ggggcgcacc cgagtcgggc catgaggccg ggaaccgcgc 180
tacaggccgt gctgctggcc gtgctgctgg tggggctgcg ggccgcgacg ggtcgcctgc 240
tgagtgcctc ggatttggac ctcagaggag ggcagccagt ctgccgggga gggacacaga 300
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<211> 382

<212> PRT

<213> Homo sapiens

<400> 137

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Asp Leu Arg Gly Gln Gln Pro Val Cys Arg Gly Gly Thr Gln Arg Pro 35 40 45

Cys Tyr Lys Val Ile Tyr Phe His Asp Thr Ser Arg Arg Leu Asn Phe 50 55 60

Glu Glu Ala Lys Glu Ala Cys Arg Arg Asp Gly Gly Gln Leu Val Ser
65 70 75 80

Ile Glu Ser Glu Asp Glu Gln Lys Leu Ile Glu Lys Phe Ile Glu Asn 85 90 95

Leu Leu Pro Ser Asp Gly Asp Phe Trp Ile Gly Leu Arg Arg Glu
100 105 110

- Glu Lys Gln Ser Asn Ser Thr Ala Cys Gln Asp Leu Tyr Ala Trp Thr 115 120 125
- Asp Gly Ser Ile Ser Gln Phe Arg Asn Trp Tyr Val Asp Glu Pro Ser 130 135
- Cys Gly Ser Glu Val Cys Val Val Met Tyr His Gln Pro Ser Ala Pro 145 150 155 160
- Ala Gly Ile Gly Gly Pro Tyr Met Phe Gln Trp Asn Asp Asp Arg Cys 165 170 175
- Asn Met Lys Asn Asn Phe Ile Cys Lys Tyr Ser Asp Glu Lys Pro Ala 180 185 190
- Val Pro Ser Arg Glu Ala Glu Gly Glu Glu Thr Glu Leu Thr Thr Pro 195 200 205
- Val Leu Pro Glu Glu Thr Gln Glu Glu Asp Ala Lys Lys Thr Phe Lys 210 215 220
- Glu Ser Arg Glu Ala Ala Leu Asn Leu Ala Tyr Ile Leu Ile Pro Ser 225 230 235 240
- Ile Pro Leu Leu Leu Leu Val Val Thr Thr Val Val Cys Trp Val 245 250 250
- Trp Ile Cys Arg Lys Arg Lys Arg Glu Gln Pro Asp Pro Ser Thr Lys 260 265 270
- Lys Gln His Thr Ile Trp Pro Ser Pro His Gln Gly Asn Ser Pro Asp 275 280 285
- Leu Glu Val Tyr Asn Val Ile Arg Lys Gln Ser Glu Ala Asp Leu Ala 290 295 300
- Glu Thr Arg Pro Asp Leu Lys Asn Ile Ser Phe Arg Val Cys Ser Gly 305 310
- Glu Ala Thr Pro Asp Asp Met Ser Cys Asp Tyr Asp Asn Met Ala Val 325 330 330
- Asn Pro Ser Glu Ser Gly Phe Val Thr Leu Val Ser Val Glu Ser Gly 340 345
- Phe Val Thr Asn Asp Ile Tyr Glu Phe Ser Pro Asp Gln Met Gly Arg 355 360 365
- Ser Lys Glu Ser Gly Trp Val Glu Asn Glu Ile Tyr Gly Tyr 370 375 380

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<210> 139
<211> 24
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      oligonucleotide probe
<400> 139
                                                                   24
aagccaaaga agcctgcagg aggg
<210> 140
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<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 140
                                                                   24
cagtccaagc ataaaggtcc tggc
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<212> DNA
<213> Homo sapiens
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ttgtgtttgc ctcctgcagc ctcaacccgg agggcagcga gggcctacca ccatgatcac 180
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cgaccgcagc ctgctgaagt tgaaaatggt gcaggtcgtg tttcgacacg gggctcggag 360
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cactaacatt tttcggaatc tggagtccac ccgttgtttg ctggctgggc ttttccagtg 720
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Lys Leu Lys Met Val Gln Val Val Phe Arg His Gly Ala Arg Ser Pro
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Leu Lys Pro Leu Pro Leu Glu Glu Gln Val Glu Trp Asn Pro Gln Leu
Leu Glu Val Pro Pro Gln Thr Gln Phe Asp Tyr Thr Val Thr Asn Leu
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Ala Gly Gly Pro Lys Pro Tyr Ser Pro Tyr Asp Ser Gln Tyr His Glu
Thr Thr Leu Lys Gly Gly Met Phe Ala Gly Gln Leu Thr Lys Val Gly
Met Gln Gln Met Phe Ala Leu Gly Glu Arg Leu Arg Lys Asn Tyr Val
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Glu Asp Ile Pro Phe Leu Ser Pro Thr Phe Asn Pro Gln Glu Val Phe
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Ile Arg Ser Thr Asn Ile Phe Arg Asn Leu Glu Ser Thr Arg Cys Leu
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Leu Ala Gly Leu Phe Gln Cys Gln Lys Glu Gly Pro Ile Ile His 180 185 190

Thr Asp Glu Ala Asp Ser Glu Val Leu Tyr Pro Asn Tyr Gln Ser Cys 195 200 205

Trp Ser Leu Arg Gln Arg Thr Arg Gly Arg Arg Gln Thr Ala Ser Leu 210 215 220

Gln Pro Gly Ile Ser Glu Asp Leu Lys Lys Val Lys Asp Arg Met Gly 225 230 235 240

Ile Asp Ser Ser Asp Lys Val Asp Phe Phe Ile Leu Leu Asp Asn Val
245 250 255

Ala Ala Glu Gln Ala His Asn Leu Pro Ser Cys Pro Met Leu Lys Arg
260 265 270

Phe Ala Arg Met Ile Glu Gln Arg Ala Val Asp Thr Ser Leu Tyr Ile 275 280 285

Leu Pro Lys Glu Asp Arg Glu Ser Leu Gln Met Ala Val Gly Pro Phe 290 295 300

Leu His Ile Leu Glu Ser Asn Leu Leu Lys Ala Met Asp Ser Ala Thr 305 310 315 320

Ala Pro Asp Lys Ile Arg Lys Leu Tyr Leu Tyr Ala Ala His Asp Val 325 330 335

Thr Phe Ile Pro Leu Leu Met Thr Leu Gly Ile Phe Asp His Lys Trp 340 345 350

Pro Pro Phe Ala Val Asp Leu Thr Met Glu Leu Tyr Gln His Leu Glu 355 360 365

Ser Lys Glu Trp Phe Val Gln Leu Tyr Tyr His Gly Lys Glu Gln Val 370 380

Pro Arg Gly Cys Pro Asp Gly Leu Cys Pro Leu Asp Met Phe Leu Asn 385 390 395 400

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<210> 143

<211> 24

<212> DNA

<213> Artificial Sequence

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| <400> 143   |     |
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| <210> 144   |     |
| <211> 24  |     |
| <212> DNA   |     |
| <213> Artificial Sequence   |     |
| <220>   |     |
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| oligonucleotide probe   |     |
| <400> 144   |     |
| gcagctctat taccacggga agga  | 24  |
| <210> 145   |     |
| <211> 24  |     |
| <212> DNA   |     |
| <213> Artificial Sequence   |     |
| <220>   |     |
| <223> Description of Artificial Sequence: Synthetic               |     |
| oligonucleotide probe   |     |
| <400> 145   |     |
| tccttcccgt ggtaatagag ctgc  | 24  |
| <210> 146   |     |
| <211> 45  |     |
| <212> DNA   |     |
| <213> Artificial Sequence   |     |
| <220>   |     |
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| oligonucleotide probe   |     |
| <400> 146   |     |
| ggcagagaac cagaggccgg aggagactgc ctctttacag ccagg                 | 45  |
| <210> 147   |     |
| <211> 1686  |     |
| <212> DNA   |     |
| <213> Homo sapiens  |     |
| <400> 147   |     |
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| cttaaatttc agctcatcac cttcacctgc cttggtcatg gctctgctat tctccttgat |     |
| cettgecatt tgeaccagae etggatteet agegteteea tetggagtge ggetggtggg | T8( |

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gagetettte teeceagtee cagagggtgt caggetgget gaeggeeetg ggeattgeaa 540
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<213> Homo sapiens
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Phe Leu Ala Ser Pro Ser Gly Val Arg Leu Val Gly Gly Leu His Arg
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Cys Asp Asp Gly Trp Asp Ile Lys Asp Val Ala Val Leu Cys Arg Glu
                        55
Leu Gly Cys Gly Ala Ala Ser Gly Thr Pro Ser Gly Ile Leu Tyr Glu
 65
                    70
                                        75
Pro Pro Ala Glu Lys Glu Gln Lys Val Leu Ile Gln Ser Val Ser Cys
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Thr Gly Thr Glu Asp Thr Leu Ala Gln Cys Glu Gln Glu Glu Val Tyr

105

100

Asp Cys Ser His Asp Glu Asp Ala Gly Ala Ser Cys Glu Asn Pro Glu 115 120 125

Ser Ser Phe Ser Pro Val Pro Glu Gly Val Arg Leu Ala Asp Gly Pro 130 135 140

Gly His Cys Lys Gly Arg Val Glu Val Lys His Gln Asn Gln Trp Tyr 145 150 155 160

Thr Val Cys Gln Thr Gly Trp Ser Leu Arg Ala Ala Lys Val Val Cys 165 170 175

Arg Gln Leu Gly Cys Gly Arg Ala Val Leu Thr Gln Lys Arg Cys Asn 180 185 190

Lys His Ala Tyr Gly Arg Lys Pro Ile Trp Leu Ser Gln Met Ser Cys 195 200 205

Ser Gly Arg Glu Ala Thr Leu Gln Asp Cys Pro Ser Gly Pro Trp Gly 210 215 220

Lys Asn Thr Cys Asn His Asp Glu Asp Thr Trp Val Glu Cys Glu Asp 225 230 235 240

Pro Phe Asp Leu Arg Leu Val Gly Gly Asp Asn Leu Cys Ser Gly Arg 245 250 255

Leu Glu Val Leu His Lys Gly Val Trp Gly Ser Val Cys Asp Asp Asn 260 265 270

Trp Gly Glu Lys Glu Asp Gln Val Val Cys Lys Gln Leu Gly Cys Gly 275 280 285

Lys Ser Leu Ser Pro Ser Phe Arg Asp Arg Lys Cys Tyr Gly Pro Gly 290 295 300

Val Gly Arg Ile Trp Leu Asp Asn Val Arg Cys Ser Gly Glu Glu Gln 305 310 315 320

Ser Leu Glu Gln Cys Gln His Arg Phe Trp Gly Phe His Asp Cys Thr 325 330 335

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<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

## oligonucleotide probe

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geteateaga gaaettaeeg etteteatge eaceaaggtg eagacacaea ageettaett 420
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1427

agggccaggg cagagaagca gcactcttag gcttgcttac tctacaaggg acagttgcat 1140 ttqttqaqac tttaatggaq atttqtctca caagtqqqaa agactgaaga aacacatctc 1200 gtgcagatct gctggcagag gacaatcaaa aacgacaaca agcttcttcc cagggtgagg 1260 ggaaacactt aaggaataaa tatggagctg gggtttaaca ctaaaaacta gaaataaaca 1320 tctcaaacag taaaaaaaaa aaaaaagggc ggccgcgact ctagagtcga cctgcagaag 1380 cttggccgcc atggcccaac ttgtttattg cagcttataa tggttac <210> 153 <211> 310 <212> PRT <213> Homo sapiens <400> 153 Met Asp Phe Ile Thr Ser Thr Ala Ile Leu Pro Leu Leu Phe Gly Cys 10 15 Leu Gly Val Phe Gly Leu Phe Arg Leu Leu Gln Trp Val Arg Gly Lys Ala Tyr Leu Arg Asn Ala Val Val Val Ile Thr Gly Ala Thr Ser Gly Leu Gly Lys Glu Cys Ala Lys Val Phe Tyr Ala Ala Gly Ala Lys Leu Val Leu Cys Gly Arg Asn Gly Gly Ala Leu Glu Leu Ile Arg Glu 70 75 Leu Thr Ala Ser His Ala Thr Lys Val Gln Thr His Lys Pro Tyr Leu Val Thr Phe Asp Leu Thr Asp Ser Gly Ala Ile Val Ala Ala Ala Ala 105 Glu Ile Leu Gln Cys Phe Gly Tyr Val Asp Ile Leu Val Asn Asn Ala Gly Ile Ser Tyr Arg Gly Thr Ile Met Asp Thr Thr Val Asp Val Asp Lys Arg Val Met Glu Thr Asn Tyr Phe Gly Pro Val Ala Leu Thr Lys 155 Ala Leu Leu Pro Ser Met Ile Lys Arg Arg Gln Gly His Ile Val Ala 165 170 Ile Ser Ser Ile Gln Gly Lys Met Ser Ile Pro Phe Arg Ser Ala Tyr 185 Ala Ala Ser Lys His Ala Thr Gln Ala Phe Phe Asp Cys Leu Arg Ala 195 200 205

Glu Met Glu Gln Tyr Glu Ile Glu Val Thr Val Ile Ser Pro Gly Tyr

|   | 210   |            |            |            |            | 215        |            |            |            |            | 220        |            |            |            |            |  |
|---|---|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|--|
| Ile<br>225  | His   | Thr        | Asn        | Leu        | Ser<br>230 | Val        | Asn        | Ala        | Ile        | Thr<br>235 | Ala        | Asp        | Gly        | Ser        | Arg<br>240 |  |
| Tyr   | Gly   | Val        | Met        | Asp<br>245 | Thr        | Thr        | Thr        | Ala        | Gln<br>250 | Gly        | Arg        | Ser        | Pro        | Val<br>255 | Glu        |  |
| Val   | Ala   | Gln        | Asp<br>260 | Val        | Leu        | Ala        | Ala        | Val<br>265 | Gly        | Lys        | Lys        | Lys        | Lys<br>270 | Asp        | Val        |  |
| Ile   | Leu   | Ala<br>275 | Asp        | Leu        | Leu        | Pro        | Ser<br>280 | Leu        | Ala        | Val        | Tyr        | Leu<br>285 | Arg        | Thr        | Leu        |  |
| Ala   | Pro<br>290  | Gly        | Leu        | Phe        | Phe        | Ser<br>295 | Leu        | Met        | Ala        | Ser        | Arg<br>300 | Ala        | Arg        | Lys        | Glu        |  |
| Arg<br>305  | Lys   | Ser        | Lys        | Asn        | Ser<br>310 |            |            |            |            |            |            |            |            |            |            |  |
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| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |   |            |            |            |            |            |            |            |            |            |            |            |            |            |            |  |
|   | <400> 154 ggtgctaaac tggtgctctg tggc 24                         |            |            |            |            |            |            |            |            |            |            |            |            |            |            |  |
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| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |   |            |            |            |            |            |            |            |            |            |            |            |            |            |            |  |
| <400> 155 cagggcaaga tgagcattcc 20  |   |            |            |            |            |            |            |            |            |            |            |            |            |            |            |  |
| <213<br><213  | 10> 156<br>11> 24<br>12> DNA<br>13> Artificial Sequence         |            |            |            |            |            |            |            |            |            |            |            |            |            |            |  |
| <220<br><220  | 3 > De  |            | iptio      |            |            |            | cial       | Seq        | uence      | e: S       | ynth       | etic       |            |            |            |  |

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<211> 50
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<212> DNA
<213> Homo sapiens
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<212> PRT

<213> Homo sapiens

<400> 159

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Gly Arg Leu Thr Ala Tyr Glu Phe Ala Lys Leu Lys Ser Lys Leu Val
50 55 60

Leu Trp Asp Ile Asn Lys His Gly Leu Glu Glu Thr Ala Ala Lys Cys
65 70 75 80

Lys Gly Leu Gly Ala Lys Val His Thr Phe Val Val Asp Cys Ser Asn 85 90 95

Arg Glu Asp Ile Tyr Ser Ser Ala Lys Lys Val Lys Ala Glu Ile Gly 100 105 110

Asp Val Ser Ile Leu Val Asn Asn Ala Gly Val Val Tyr Thr Ser Asp 115 120 125

Leu Phe Ala Thr Gln Asp Pro Gln Ile Glu Lys Thr Phe Glu Val Asn 130 135 140

Val Leu Ala His Phe Trp Thr Thr Lys Ala Phe Leu Pro Ala Met Thr 145 150 155 160

Lys Asn Asn His Gly His Ile Val Thr Val Ala Ser Ala Ala Gly His 165 170 175

Val Ser Val Pro Phe Leu Leu Ala Tyr Cys Ser Ser Lys Phe Ala Ala 180 185 190

Val Gly Phe His Lys Thr Leu Thr Asp Glu Leu Ala Ala Leu Gln Ile 195 200 205

Thr Gly Val Lys Thr Thr Cys Leu Cys Pro Asn Phe Val Asn Thr Gly 210 215 220

Phe Ile Lys Asn Pro Ser Thr Ser Leu Gly Pro Thr Leu Glu Pro Glu 225 230 235 240

Glu Val Val Asn Arg Leu Met His Gly Ile Leu Thr Glu Gln Lys Met 245 250 255

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<210> 162
<211> 48
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Ile Glu Ala Gly Lys Ile Gln Lys Gly Arg Glu Leu Ser Leu Val Gly

Pro Phe Pro Gly Leu Asn Met Lys Ser Tyr Ala Gly Phe Leu Thr Val

|            |            |            |            |            |            |            |            |            |            |            |            |            |            |            | 80         |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
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| Ile        | Gln        | Pro        | Glu<br>100 | Asp        | Ala        | Pro        | Val        | Val<br>105 | Leu        | Trp        | Leu        | Gln        | Gly<br>110 | Gly        | Pro        |
| Gly        | Gly        | Ser<br>115 | Ser        | Met        | Phe        | Gly        | Leu<br>120 | Phe        | Val        | Glu        | His        | Gly<br>125 | Pro        | Tyr        | Val        |
| Val        | Thr<br>130 | Ser        | Asn        | Met        | Thr        | Leu<br>135 | Arg        | Asp        | Arg        | Asp        | Phe<br>140 | Pro        | Trp        | Thr        | Thr        |
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| Tyr        | Lys        | Asn<br>195 | Asn        | Asp        | Phe        | Tyr        | Val<br>200 | Thr        | Gly        | Glu        | Ser        | Tyr<br>205 | Ala        | Gly        | Lys        |
| Tyr        | Val<br>210 | Pro        | Ala        | Ile        | Ala        | His<br>215 | Leu        | Ile        | His        | Ser        | Leu<br>220 | Asn        | Pro        | Val        | Arg        |
| Glu<br>225 | Val        | Lys        | Ile        | Asn        | Leu<br>230 | Asn        | Gly        | Ile        | Ala        | Ile<br>235 | Gly        | Asp        | Gly        | Tyr        | Ser<br>240 |
| Asp        | Pro        | Glu        | Ser        | Ile<br>245 | Ile        | Gly        | Gly        | Tyr        | Ala<br>250 | Glu        | Phe        | Leu        | Tyr        | Gln<br>255 | Ile        |
| Gly        | Leu        | Leu        | Asp<br>260 | Glu        | Lys        | Gln        | Lys        | Lys<br>265 | Tyr        | Phe        | Gln        | Lys        | Gln<br>270 | Cys        | His        |
| Glu        |            | Ile<br>275 |            | His        | Ile        | Arg        | Lys<br>280 | Gln        | Asn        | Trp        | Phe        | Glu<br>285 |            | Phe        | Glu        |
| Ile        | Leu<br>290 | Asp        | Lys        | Leu        | Leu        | Asp<br>295 | Gly        | Asp        | Leu        | Thr        | Ser<br>300 | Asp        | Pro        | Ser        | Туг        |
| Phe<br>305 | Gln        | Asn        | Val        | Thr        | Gly<br>310 | Cys        | Ser        | Asn        | Tyr        | Tyr<br>315 | Asn        | Phe        | Leu        | Arg        | Cys<br>320 |
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| Glu        | Val        | Arg        | Gln        |            | Ile        | His        | Val        | Gly        |            | Gln        | Thr        | Phe        | Asn        |            | Gl         |

| Thr Ile Val Glu Lys Tyr Leu Arg Glu Asp Thr Val Gln Ser Val Lys 365             |    |  |  |  |  |  |  |  |  |  |  |
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| Gly Gln Leu Asp Ile Ile Val Ala Ala Ala Leu Thr Glu Arg Ser Leu 385 390 400     |    |  |  |  |  |  |  |  |  |  |  |
| Met Gly Met Asp Trp Lys Gly Ser Gln Glu Tyr Lys Lys Ala Glu Lys 405 410 415     |    |  |  |  |  |  |  |  |  |  |  |
| Lys Val Trp Lys Ile Phe Lys Ser Asp Ser Glu Val Ala Gly Tyr Ile<br>420 425 430  |    |  |  |  |  |  |  |  |  |  |  |
| Arg Gln Ala Gly Asp Phe His Gln Val Ile Ile Arg Gly Gly His 435 440 445         |    |  |  |  |  |  |  |  |  |  |  |
| Ile Leu Pro Tyr Asp Gln Pro Leu Arg Ala Phe Asp Met Ile Asn Arg<br>450 455 460  | ;  |  |  |  |  |  |  |  |  |  |  |
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Leu Leu Ala Pro Pro Ala Ala Gly Met Pro Gln Phe Ser Thr Phe His
Ser Glu Asn Arg Asp Trp Thr Phe Asn His Leu Thr Val His Gln Gly
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            100
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Gly Asn Leu Thr Ile Gln Val Ala His Lys Thr Gly Pro Glu Glu Asp
                       135
    130
Asn Lys Ser Arg Tyr Pro Pro Leu Ile Val Gln Pro Cys Ser Glu Val
145
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                                       155
Leu Thr Leu Thr Asn Asn Val Asn Lys Leu Leu Ile Ile Asp Tyr Ser
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                                   170
                165
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His His Pro Pro Asp Asp Ser Ala Leu Cys Ala Phe Pro Ile Arg Ala

Ile Asn Leu Gln Ile Lys Glu Arg Leu Gln Ser Cys Tyr Gln Gly Glu
420 425 430

Gly Asn Leu Glu Leu Asn Trp Leu Leu Gly Lys Asp Val Gln Cys Thr
435 440 445

Lys Ala Pro Val Pro Ile Asp Asp Asn Phe Cys Gly Leu Asp Ile Asn

410

390

|   | 450  |            |            |            |                | 455        |            |            |            |            | 460        |            |            |            |            |  |
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| Gln<br>465  | Pro  | Leu        | Gly        | Gly        | Ser<br>470     | Thr        | Pro        | Val        | Glu        | Gly<br>475 | Leu        | Thr        | Leu        | Tyr        | Thr<br>480 |  |
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| Tyr   | Ser  | Val        | Val<br>500 | Phe        | Val            | Gly        | Thr        | Lys<br>505 | Ser        | Gly        | Lys        | Leu        | Lys<br>510 | Lys        | Val        |  |
| Arg   | Val  | Tyr<br>515 | Glu        | Phe        | Arg            | Cys        | Ser<br>520 | Asn        | Ala        | Ile        | His        | Leu<br>525 | Leu        | Ser        | Lys        |  |
| Glu   | Ser<br>530   | Leu        | Leu        | Glu        | Gly            | Ser<br>535 | Tyr        | Trp        | Trp        | Arg        | Phe<br>540 | Asn        | Tyr        | Arg        | Gln        |  |
| Leu<br>545  | Tyr  | Phe        | Leu        | Gly        | Glu<br>550     | Gln        | Arg        |            |            |            |            |            |            |            |            |  |
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130

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Ser Thr Leu Val Pro Leu Arg Leu Arg His Arg Gln Leu Gly Leu Gln
                                                 45
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Ala Lys Gly Trp Asn Phe Met Leu Glu Asp Ser Thr Phe Trp Ile Phe
Gly Gly Ser Ile His Tyr Phe Arg Val Pro Arg Glu Tyr Trp Arg Asp
                     70
                                         75
Arg Leu Leu Lys Met Lys Ala Cys Gly Leu Asn Thr Leu Thr Thr Tyr
                 85
Val Pro Trp Asn Leu His Glu Pro Glu Arq Gly Lys Phe Asp Phe Ser
                                105
Gly Asn Leu Asp Leu Glu Ala Phe Val Leu Met Ala Ala Glu Ile Gly
        115
Leu Trp Val Ile Leu Arg Pro Gly Pro Tyr Ile Cys Ser Glu Met Asp
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135

Leu Gly Gly Leu Pro Ser Trp Leu Leu Gln Asp Pro Gly Met Arg Leu 145 Arg Thr Thr Tyr Lys Gly Phe Thr Glu Ala Val Asp Leu Tyr Phe Asp 165 His Leu Met Ser Arg Val Val Pro Leu Gln Tyr Lys Arg Gly Gly Pro 185 Ile Ile Ala Val Gln Val Glu Asn Glu Tyr Gly Ser Tyr Asn Lys Asp Pro Ala Tyr Met Pro Tyr Val Lys Lys Ala Leu Glu Asp Arg Gly Ile 215 Val Glu Leu Leu Thr Ser Asp Asn Lys Asp Gly Leu Ser Lys Gly 235 Ile Val Gln Gly Val Leu Ala Thr Ile Asn Leu Gln Ser Thr His Glu 250 Leu Gln Leu Leu Thr Thr Phe Leu Phe Asn Val Gln Gly Thr Gln Pro 265 Lys Met Val Met Glu Tyr Trp Thr Gly Trp Phe Asp Ser Trp Gly Gly 275 Pro His Asn Ile Leu Asp Ser Ser Glu Val Leu Lys Thr Val Ser Ala 295 Ile Val Asp Ala Gly Ser Ser Ile Asn Leu Tyr Met Phe His Gly Gly 305 310 315 Thr Asn Phe Gly Phe Met Asn Gly Ala Met His Phe His Asp Tyr Lys 330 Ser Asp Val Thr Ser Tyr Asp Tyr Asp Ala Val Leu Thr Glu Ala Gly 345 Asp Tyr Thr Ala Lys Tyr Met Lys Leu Arg Asp Phe Phe Gly Ser Ile 355 Ser Gly Ile Pro Leu Pro Pro Pro Pro Asp Leu Leu Pro Lys Met Pro 375 Tyr Glu Pro Leu Thr Pro Val Leu Tyr Leu Ser Leu Trp Asp Ala Leu 395 Lys Tyr Leu Gly Glu Pro Ile Lys Ser Glu Lys Pro Ile Asn Met Glu 410 405 Asn Leu Pro Val Asn Gly Gly Asn Gly Gln Ser Phe Gly Tyr Ile Leu

425

Tyr Glu Thr Ser Ile Thr Ser Ser Gly Ile Leu Ser Gly His Val His 440 435 Asp Arg Gly Gln Val Phe Val Asn Thr Val Ser Ile Gly Phe Leu Asp 455 Tyr Lys Thr Thr Lys Ile Ala Val Pro Leu Ile Gln Gly Tyr Thr Val 465 470 475 480 Leu Arg Ile Leu Val Glu Asn Arg Gly Arg Val Asn Tyr Gly Glu Asn 485 Ile Asp Asp Gln Arg Lys Gly Leu Ile Gly Asn Leu Tyr Leu Asn Asp 505 Ser Pro Leu Lys Asn Phe Arg Ile Tyr Ser Leu Asp Met Lys Lys Ser 520 515 Phe Phe Gln Arg Phe Gly Leu Asp Lys Trp Xaa Ser Leu Pro Glu Thr 535 Pro Thr Leu Pro Ala Phe Phe Leu Gly Ser Leu Ser Ile Ser Ser Thr 550 545 555 Pro Cys Asp Thr Phe Leu Lys Leu Glu Gly Trp Glu Lys Gly Val Val Phe Ile Asn Gly Gln Asn Leu Gly Arg Tyr Trp Asn Ile Gly Pro Gln 585 590 Lys Thr Leu Tyr Leu Pro Gly Pro Trp Leu Ser Ser Gly Ile Asn Gln Val Ile Val Phe Glu Glu Thr Met Ala Gly Pro Ala Leu Gln Phe Thr 615 Glu Thr Pro His Leu Gly Arg Asn Gln Tyr Ile Lys 625 630 <210> 176 <211> 2505 <212> DNA <213> Homo sapiens <400> 176 ggggacgcgg agctgagagg ctccgggcta gctaggtgta gggtggacg ggtcccagga 60 ccctggtgag ggttctctac ttggccttcg gtgggggtca agacgcaggc acctacgcca 120 aaggggagca aagccgggct cggcccgagg cccccaggac ctccatctcc caatgttgga 180 ggaatccgac acgtgacggt ctgtccgccg tctcagacta gaggagcgct gtaaacgcca 240 tggctcccaa gaagctgtcc tgccttcgtt ccctgctgct gccgctcagc ctgacgctac 300

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<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

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Val Ser Gly Ser Leu His Tyr Phe Arg Val Pro Arg Val Leu Trp Ala

- Asp Arg Leu Leu Lys Met Arg Trp Ser Gly Leu Asn Ala Ile Gln Phe 65 70 75 80
- Tyr Val Pro Trp Asn Tyr His Glu Pro Gln Pro Gly Val Tyr Asn Phe 85 90 95
- Asn Gly Ser Arg Asp Leu Ile Ala Phe Leu Asn Glu Ala Ala Leu Ala 100 105 110
- Asn Leu Leu Val Ile Leu Arg Pro Gly Pro Tyr Ile Cys Ala Glu Trp 115 120 125
- Glu Met Gly Gly Leu Pro Ser Trp Leu Leu Arg Lys Pro Glu Ile His 130 135 140
- Leu Arg Thr Ser Asp Pro Asp Phe Leu Ala Ala Val Asp Ser Trp Phe 145 150 155 160
- Lys Val Leu Leu Pro Lys Ile Tyr Pro Trp Leu Tyr His Asn Gly Gly
  165 170 175
- Asn Ile Ile Ser Ile Gln Val Glu Asn Glu Tyr Gly Ser Tyr Arg Ala 180 185 190
- Cys Asp Phe Ser Tyr Met Arg His Leu Ala Gly Leu Phe Arg Ala Leu 195 200 205
- Leu Gly Glu Lys Ile Leu Leu Phe Thr Thr Asp Gly Pro Glu Gly Leu 210 215 220
- Lys Cys Gly Ser Leu Arg Gly Leu Tyr Thr Thr Val Asp Phe Gly Pro 225 230 235
- Ala Asp Asn Met Thr Lys Ile Phe Thr Leu Leu Arg Lys Tyr Glu Pro 245 250 255
- His Gly Pro Leu Val Asn Ser Glu Tyr Tyr Thr Gly Trp Leu Asp Tyr 260 265 270
- Trp Gly Gln Asn His Ser Thr Arg Ser Val Ser Ala Val Thr Lys Gly 275 280 285
- Leu Glu Asn Met Leu Lys Leu Gly Ala Ser Val Asn Met Tyr Met Phe 290 295 300
- His Gly Gly Thr Asn Phe Gly Tyr Trp Asn Gly Ala Asp Lys Lys Gly 305 310 315
- Arg Phe Leu Pro Ile Thr Thr Ser Tyr Asp Tyr Asp Ala Pro Ile Ser 325 330 335
- Glu Ala Gly Asp Pro Thr Pro Lys Leu Phe Ala Leu Arg Asp Val Ile

Leu Phe Pro Arg Gly Ala Leu Asn Lys Ile Thr Leu Leu Glu Leu Glu 

Asp Val Pro Leu Gln Pro Gln Val Gln Phe Leu Asp Lys Pro Ile Leu

| Asn Ser Thr Ser Thr Leu His Arg Thr His Ile Asn Ser Leu Ser Ala 625 630 635 640 |    |
|---|----|
| Asp Thr Leu Ser Ala Ser Glu Pro Met Glu Leu Ser Gly His 645 650                 |    |
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| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |    |
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| <210> 180<br><211> 50<br><212> DNA<br><213> Artificial Sequence                 |    |
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| <400> 180 gggcttcacc gaagcagtgg acctttattt tgaccacctg atgtccaggg                | 50 |
| <210> 181<br><211> 22<br><212> DNA<br><213> Artificial Sequence                 |    |
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gtgtttatgg ctttatctgc ctctacactc tcttctggtt attcaggata cctttgaagg 180
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1947

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Phe Ser Asp Ile Pro Asp Val Lys Asn Asp Phe Ala Phe Leu Leu His
Met Val Asp Gln Tyr Asp Gln Leu Tyr Ser Lys Arg Phe Gly Val Phe
 65
                     70
                                         75
Leu Ser Glu Val Ser Glu Asn Lys Leu Arg Glu Ile Ser Leu Asn His
Glu Trp Thr Phe Glu Lys Leu Arq Gln His Ile Ser Arg Asn Ala Gln
                                105
Asp Lys Gln Glu Leu His Leu Phe Met Leu Ser Gly Val Pro Asp Ala
        115
                            120
Val Phe Asp Leu Thr Asp Leu Asp Val Leu Lys Leu Glu Leu Ile Pro
                        135
Glu Ala Lys Ile Pro Ala Lys Ile Ser Gln Met Thr Asn Leu Gln Glu
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145
                    150
                                        155
Leu His Leu Cys His Cys Pro Ala Lys Val Glu Gln Thr Ala Phe Ser
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Phe Leu Arg Asp His Leu Arg Cys Leu His Val Lys Phe Thr Asp Val

185

180

- Ala Glu Ile Pro Ala Trp Val Tyr Leu Leu Lys Asn Leu Arg Glu Leu 195 200 205
- Tyr Leu Ile Gly Asn Leu Asn Ser Glu Asn Asn Lys Met Ile Gly Leu 210 215 220
- Glu Ser Leu Arg Glu Leu Arg His Leu Lys Ile Leu His Val Lys Ser 225 230 235 240
- Asn Leu Thr Lys Val Pro Ser Asn Ile Thr Asp Val Ala Pro His Leu 245 250 255
- Thr Lys Leu Val Ile His Asn Asp Gly Thr Lys Leu Leu Val Leu Asn 260 265 270
- Ser Leu Lys Lys Met Met Asn Val Ala Glu Leu Glu Leu Gln Asn Cys 275 280 285
- Glu Leu Glu Arg Ile Pro His Ala Ile Phe Ser Leu Ser Asn Leu Gln 290 295 300
- Glu Leu Asp Leu Lys Ser Asn Asn Ile Arg Thr Ile Glu Glu Ile Ile 305 310 315 320
- Ser Phe Gln His Leu Lys Arg Leu Thr Cys Leu Lys Leu Trp His Asn 325 330 335
- Lys Ile Val Thr Ile Pro Pro Ser Ile Thr His Val Lys Asn Leu Glu 340 345 350
- Ser Leu Tyr Phe Ser Asn Asn Lys Leu Glu Ser Leu Pro Val Ala Val 355 360 365
- Phe Ser Leu Gln Lys Leu Arg Cys Leu Asp Val Ser Tyr Asn Asn Ile 370 . 375 380
- Ser Met Ile Pro Ile Glu Ile Gly Leu Leu Gln Asn Leu Gln His Leu 385 390 395 400
- His Ile Thr Gly Asn Lys Val Asp Ile Leu Pro Lys Gln Leu Phe Lys 405 410 415
- Cys Ile Lys Leu Arg Thr Leu Asn Leu Gly Gln Asn Cys Ile Thr Ser 420 425 430
- Leu Pro Glu Lys Val Gly Gln Leu Ser Gln Leu Thr Gln Leu Glu Leu 435 440 445
- Lys Gly Asn Cys Leu Asp Arg Leu Pro Ala Gln Leu Gly Gln Cys Arg
  450 455 460
- Met Leu Lys Lys Ser Gly Leu Val Val Glu Asp His Leu Phe Asp Thr

| 465   | 470                        | 475   | 480              |  |  |  |  |  |  |  |  |
|---|----------------------------|---|------------------|--|--|--|--|--|--|--|--|
| Leu Pro Leu Glu   | ı Val Lys Glu Ala :<br>485 | Leu Asn Gln Asp Ile Ası<br>490  | n Ile Pro<br>495 |  |  |  |  |  |  |  |  |
| Phe Ala Asn Gly Ile<br>500  |                            |   |                  |  |  |  |  |  |  |  |  |
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| <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe |                            |   |                  |  |  |  |  |  |  |  |  |
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| <210> 187<br><211> 24<br><212> DNA<br><213> Artificia                           | al Sequence                |   |                  |  |  |  |  |  |  |  |  |
| _   | on of Artificial a         | Sequence: Synthetic   |                  |  |  |  |  |  |  |  |  |
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| <del>-</del>  | eotide probe               | sequence. Synthetic   |                  |  |  |  |  |  |  |  |  |
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| <210> 189<br><211> 2917<br><212> DNA<br><213> Homo sapi                         | .ens                       |   |                  |  |  |  |  |  |  |  |  |
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- Thr Val Ser Leu Gly Gly Ala Asn Met Ala Glu Thr His Lys Ala Met
- Ile Leu Gln Leu Asn Pro Ser Glu Asn Cys Thr Trp Thr Ile Glu Arg
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- Pro Glu Asn Lys Ser Ile Arg Ile Ile Phe Ser Tyr Val Gln Leu Asp 65 70 75 80
- Pro Asp Gly Ser Cys Glu Ser Glu Asn Ile Lys Val Phe Asp Gly Thr 85 90 95
- Ser Ser Asn Gly Pro Leu Leu Gly Gln Val Cys Ser Lys Asn Asp Tyr 100 105 110
- Val Pro Val Phe Glu Ser Ser Ser Ser Thr Leu Thr Phe Gln Ile Val 115 120 125
- Thr Asp Ser Ala Arg Ile Gln Arg Thr Val Phe Val Phe Tyr Tyr Phe 130 135 140
- Phe Ser Pro Asn Ile Ser Ile Pro Asn Cys Gly Gly Tyr Leu Asp Thr 145 150 155 160
- Leu Glu Gly Ser Phe Thr Ser Pro Asn Tyr Pro Lys Pro His Pro Glu 165 170 175
- Leu Ala Tyr Cys Val Trp His Ile Gln Val Glu Lys Asp Tyr Lys Ile 180 185 190
- Lys Leu Asn Phe Lys Glu Ile Phe Leu Glu Ile Asp Lys Gln Cys Lys
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- Ile Gly Gln Val Cys Gly Arg Val Thr Pro Thr Phe Glu Ser Ser Ser 225 230 235 240
- Asn Ser Leu Thr Val Val Leu Ser Thr Asp Tyr Ala Asn Ser Tyr Arg 245 250 255
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- Thr Ser Leu Thr Cys Ser Ser Asp Arg Met Arg Val Ile Ile Ser Lys 275 280 285

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- Asp Pro Thr Cys Arg Pro Lys Leu Ser Asn Val Val Glu Phe Ser Val 305 310 315
- Pro Leu Asn Gly Cys Gly Thr Ile Arg Lys Val Glu Asp Gln Ser Ile 325 330 335
- Thr Tyr Thr Asn Ile Ile Thr Phe Ser Ala Ser Ser Thr Ser Glu Val
- Ile Thr Arg Gln Lys Gln Leu Gln Ile Ile Val Lys Cys Glu Met Gly 355 360 365
- His Asn Ser Thr Val Glu Ile Ile Tyr Ile Thr Glu Asp Asp Val Ile 370 375 380
- Gln Ser Gln Asn Ala Leu Gly Lys Tyr Asn Thr Ser Met Ala Leu Phe 385 390 395 400
- Glu Ser Asn Ser Phe Glu Lys Thr Ile Leu Glu Ser Pro Tyr Tyr Val 405 410 415
- Asp Leu Asn Gln Thr Leu Phe Val Gln Val Ser Leu His Thr Ser Asp 420 425 430
- Pro Asn Leu Val Val Phe Leu Asp Thr Cys Arg Ala Ser Pro Thr Ser 435 440 445
- Asp Phe Ala Ser Pro Thr Tyr Asp Leu Ile Lys Ser Gly Cys Ser Arg 450 455 460
- Asp Glu Thr Cys Lys Val Tyr Pro Leu Phe Gly His Tyr Gly Arg Phe 465 470 475 480
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- Asn Gln Gly Cys Val Ser Arg Ser Lys Arg Asp Ile Ser Ser Tyr Lys 515 520 525
- Trp Lys Thr Asp Ser Ile Ile Gly Pro Ile Arg Leu Lys Arg Asp Arg
- Ser Ala Ser Gly Asn Ser Gly Phe Gln His Glu Thr His Ala Glu Glu 545 550 555 560
- Thr Pro Asn Gln Pro Phe Asn Ser Val His Leu Phe Ser Phe Met Val

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- Ser Val Pro Ser Phe Gly Ser Glu Trp Phe Trp Trp Tyr Trp Gln Lys
- Glu Lys Ile Pro Lys Tyr Val Glu Phe Met Lys Asp Asn Tyr Pro Pro
- Ser Phe Lys Tyr Glu Asp Phe Gly Pro Leu Phe Thr Ala Lys Phe Phe 100 105 110
- Asn Ala Asn Gln Trp Ala Asp Ile Phe Gln Ala Ser Gly Ala Lys Tyr 115 120 125
- Ile Val Leu Thr Ser Lys His His Glu Gly Phe Thr Leu Trp Gly Ser 130 135 140
- Glu Tyr Ser Trp Asn Trp Asn Ala Ile Asp Glu Gly Pro Lys Arg Asp 145 150 155 160
- Ile Val Lys Glu Leu Glu Val Ala Ile Arg Asn Arg Thr Asp Leu Arg 165 170 175
- Phe Gly Leu Tyr Tyr Ser Leu Phe Glu Trp Phe His Pro Leu Phe Leu
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- Glu Asp Glu Ser Ser Ser Phe His Lys Arg Gln Phe Pro Val Ser Lys
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- Thr Leu Pro Glu Leu Tyr Glu Leu Val Asn Asn Tyr Gln Pro Glu Val 210 215 220
- Leu Trp Ser Asp Gly Asp Gly Gly Ala Pro Asp Gln Tyr Trp Asn Ser 225 230 235 240
- Thr Gly Phe Leu Ala Trp Leu Tyr Asn Glu Ser Pro Val Arg Gly Thr 245 250 255
- Val Val Thr Asn Asp Arg Trp Gly Ala Gly Ser Ile Cys Lys His Gly 260 265 270
- Gly Phe Tyr Thr Cys Ser Asp Arg Tyr Asn Pro Gly His Leu Leu Pro
- His Lys Trp Glu Asn Cys Met Thr Ile Asp Lys Leu Ser Trp Gly Tyr 290 295 300
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| Leu   | Arg            | Gln<br>355 | Val        | Gly          | Ser        | Trp        | Leu<br>360 | Lys        | Val        | Asn        | Gly        | Glu<br>365 | Ala        | Ile        | Tyr        |    |
| Glu   | Thr<br>370     | Tyr        | Thr        | Trp          | Arg        | Ser<br>375 | Gln        | Asn        | Asp        | Thr        | Val<br>380 | Thr        | Pro        | Asp        | Val        |    |
| Trp<br>385  | Tyr            | Thr        | Ser        | Lys          | Pro<br>390 | Lys        | Glu        | Lys        | Leu        | Val<br>395 | Tyr        | Ala        | Ile        | Phe        | Leu<br>400 |    |
| Lys   | Trp            | Pro        | Thr        | Ser<br>405   | Gly        | Gln        | Leu        | Phe        | Leu<br>410 | Gly        | His        | Pro        | Lys        | Ala<br>415 | Ile        |    |
| Leu   | Gly            | Ala        | Thr<br>420 | Glu          | Val        | Lys        | Leu        | Leu<br>425 | Gly        | His        | Gly        | Gln        | Pro<br>430 | Leu        | Asn        |    |
| Tr  | Ile            | Ser<br>435 | Leu        | Glu          | Gln        | Asn        | Gly<br>440 | Ile        | Met        | Val        | Glu        | Leu<br>445 | Pro        | Gln        | Leu        |    |
| Thi   | 1le<br>450     |            | Gln        | Met          | Pro        | Cys<br>455 | Lys        | Trp        | Gly        | Trp        | Ala<br>460 | Leu        | Ala        | Leu        | Thr        |    |
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Ser Phe Glu Thr Cys Ser Tyr Gly Trp Val Gly Asp Gly Phe Val Val
                 85
Ile Ser Arq Ile Ser Pro Asn Pro Lys Cys Gly Lys Asn Gly Val Gly
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24

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<210> 204
<211> 24
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Ala Ser Glu His Pro Lys Ile Glu Phe Tyr Leu Lys Asn Phe Thr Ser 210 215 220

Ala Lys Asp Val Leu Phe Ala Ile Lys Glu Val Gly Phe Arg Gly Gly 225 230 235

Asn Ser Asn Thr Gly Lys Ala Leu Lys His Thr Ala Gln Lys Phe Phe 245 250 255

- Thr Val Asp Ala Gly Val Arg Lys Gly Ile Pro Lys Val Val Val Val 260 265 270
- Phe Ile Asp Gly Trp Pro Ser Asp Asp Ile Glu Glu Ala Gly Ile Val 275 280 285
- Ala Arg Glu Phe Gly Val Asn Val Phe Ile Val Ser Val Ala Lys Pro 290 295 300
- Ile Pro Glu Glu Leu Gly Met Val Gln Asp Val Thr Phe Val Asp Lys 305 310 315 320
- Ala Val Cys Arg Asn Asn Gly Phe Phe Ser Tyr His Met Pro Asn Trp 325 330 335
- Phe Gly Thr Thr Lys Tyr Val Lys Pro Leu Val Gln Lys Leu Cys Thr 340 345 350
- His Glu Gln Met Met Cys Ser Lys Thr Cys Tyr Asn Ser Val Asn Ile 355 360 365
- Ala Phe Leu Ile Asp Gly Ser Ser Ser Val Gly Asp Ser Asn Phe Arg 370 375 380
- Leu Met Leu Glu Phe Val Ser Asn Ile Ala Lys Thr Phe Glu Ile Ser 385 390 395 400
- Asp Ile Gly Ala Lys Ile Ala Ala Val Gln Phe Thr Tyr Asp Gln Arg
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- Thr Glu Phe Ser Phe Thr Asp Tyr Ser Thr Lys Glu Asn Val Leu Ala 420 425 430
- Val Ile Arg Asn Ile Arg Tyr Met Ser Gly Gly Thr Ala Thr Gly Asp 435 440 445
- Ala Ile Ser Phe Thr Val Arg Asn. Val Phe Gly Pro Ile Arg Glu Ser 450 455 460
- Pro Asn Lys Asn Phe Leu Val Ile Val Thr Asp Gly Gln Ser Tyr Asp 465 470 475 480
- Asp Val Gln Gly Pro Ala Ala Ala Ala His Asp Ala Gly Ile Thr Ile 485 490 495
- Phe Ser Val Gly Val Ala Trp Ala Pro Leu Asp Asp Leu Lys Asp Met 500 505 510
- Ala Ser Lys Pro Lys Glu Ser His Ala Phe Phe Thr Arg Glu Phe Thr 515 520 525
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Leu Arg Ser Ala Val Glu Glu Met Glu Ala Glu Glu Ala Ala Lys 70

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- Arg Glu Ile His Lys Ile Thr Asn Asn Gln Thr Gly Gln Met Val Phe
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- Ser Glu Thr Val Ile Thr Ser Val Gly Asp Glu Glu Gly Arg Arg Ser 130 135 140
- His Glu Cys Ile Ile Asp Glu Asp Cys Gly Pro Ser Met Tyr Cys Gln 145 150 155 160
- Phe Ala Ser Phe Gln Tyr Thr Cys Gln Pro Cys Arg Gly Gln Arg Met 165 170 175
- Leu Cys Thr Arg Asp Ser Glu Cys Cys Gly Asp Gln Leu Cys Val Trp
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- Gly His Cys Thr Lys Met Ala Thr Arg Gly Ser Asn Gly Thr Ile Cys 195 200 205
- Asp Asn Gln Arg Asp Cys Gln Pro Gly Leu Cys Cys Ala Phe Gln Arg 210 215 220
- Gly Leu Leu Phe Pro Val Cys Thr Pro Leu Pro Val Glu Gly Glu Leu 225 230 235 240
- Cys His Asp Pro Ala Ser Arg Leu Leu Asp Leu Ile Thr Trp Glu Leu 245 250 255
- Glu Pro Asp Gly Ala Leu Asp Arg Cys Pro Cys Ala Ser Gly Leu Leu 260 265 270
- Cys Gln Pro His Ser His Ser Leu Val Tyr Val Cys Lys Pro Thr Phe 275 280 285
- Val Gly Ser Arg Asp Gln Asp Gly Glu Ile Leu Leu Pro Arg Glu Val 290 295 300
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| <210><211><212><212><213> | 24   |    |
| <220><br><223>            | Synthetic Oligonucleotide Probe                            |    |
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| <210><211><212><213>      | 18   |    |
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| <210><211><211><212><213> | 18   |    |
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Ser Asp Ala Arg Asp Cys Asp Phe His Ala Leu Pro Gln Leu Leu 110 115 120

Ser Leu His Leu Glu Glu Asn Gln Leu Thr Arg Leu Glu Asp His 125 130 135

Ser Phe Ala Gly Leu Ala Ser Leu Gln Glu Leu Tyr Leu Asn His

| Asn | Gln   | Leu  | ΤŊ   |     | Arg<br>155      | Ile   | Ala   | Pro   | Arg   | Ala<br>160   | Phe        | Ser   | Gly   | Leu   | Ser<br>165   |
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| Asn | Leu   | Leu  | ı Aı |     | Leu<br>170      | His   | Leu   | Asn   | Ser   | Asn<br>175   | Leu        | Leu   | Arg   | Ala   | Ile<br>180   |
| Asp | Ser   | Arg  | y T  |     | Phe<br>185      | Glu   | Met   | Leu   | Pro   | Asn<br>190   | Leu        | Glu   | Ile   | Leu   | Met<br>195   |
| Ile | Gly   | Gly  | / A  |     | Lys<br>200      | Val   | Asp   | Ala   | Ile   | Leu<br>205   | Asp        | Met   | Asn   | Phe   | Arg<br>210   |
| Pro | Leu   | Ala  | a A  |     | Leu<br>215      | Arg   | Ser   | Leu   | Val   | Leu<br>220   | Ala        | Gly   | Met   | Asn   | Leu<br>225   |
| Arg | Glu   | Ile  | e S  | er  | Asp<br>230      | Tyr   | Ala   | Leu   | Glu   | Gly<br>235   | Leu        | Gln   | Ser   | Leu   | Glu<br>240   |
| Ser | Leu   | . Se | r P  | he  | Туr<br>245      | Asp   | Asn   | Gln   | Leu   | Ala<br>250   | Arg        | Val   | Pro   | Arg   | Arg<br>255   |
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| Asn | Pro   | Le   | u G  | ln  | Arg<br>275      |       | Gly   | Pro   | Gly   | Asp<br>280   | Phe        | Ala   | Asn   | Met   | Leu<br>285   |
| His | Leu   | ı Ly | s C  | Glu | Leu<br>290      | Gly   | Leu   | Asn   | Asn   | Met<br>295   | Glu        | Glu   | ı Lev | ı Val | Ser<br>300   |
| Il€ | e Asp | р Ьу | s I  | Phe | Ala<br>305      |       | ı Val | . Asr | Leu   | 310          | Glu        | Lev   | ı Thi | Lys   | 315          |
| Asp | , Ile | e Tł | ır A | Asn | Asn<br>320      |       | Arg   | J Let | ı Ser | 2 Phe<br>325 | e Ile      | e His | s Pro | o Arg | g Ala<br>330 |
| Phe | e His | s Hi | is 1 | Leu | Pro<br>335      |       | n Met | ; Glı | ı Thi | Let<br>340   | ı Met      | . Le  | ı Ası | n Asi | n Asn<br>345 |
| Ala | a Le  | u Se | er.  | Ala | Le:             | ı His | s Glı | n Gli | n Thi | r Va:        | l Gl:<br>5 | ı Se: | r Le  | u Pr  | o Asr<br>360 |
| Le  | u Gl  | n G  | lu   | Val | Gl <sub>3</sub> |       | ı Hi  | s Gl  | y Ası | n Pro        | o Ilo      | e Ar  | g Cy  | s As  | p Cys<br>375 |
| Va  | 1 11  | e A  | rg   | Trp | Ala<br>38       |       | n Al  | a Th  | r Gl  | y Th<br>38   | r Ar       | g Va  | l Ar  | g Ph  | e Ile<br>390 |
| Gl  | u Pr  | o G  | ln   | Ser | Th:             | r Le  | u Cy  | s Al  | a Gl  | u Pr<br>40   | o Pr<br>O  | o As  | p Le  | u Gl  | n Arg<br>40  |
| Le  | u Pr  | :o V | al   | Arc | Gl              | u Va  | l Pr  | o Ph  | e Ar  | g Gl         | u Me       | t Th  | r As  | p Hi  | s Cy         |

|     |     |     |     | 410        |     |     |      |     | 415        |     |     |     |     | 420        |
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| Ala | Ser | Gly | Glu | Ser<br>440 | Met | Val | Leu  | His | Cys<br>445 | Arg | Ala | Leu | Ala | Glu<br>450 |
| Pro | Glu | Pro | Glu | Ile<br>455 | Tyr | Trp | Val  | Thr | Pro<br>460 | Ala | Gly | Leu | Arg | Leu<br>465 |
| Thr | Pro | Ala | His | Ala<br>470 | Gly | Arg | Arg  | Tyr | Arg<br>475 | Val | Tyr | Pro | Glu | Gly<br>480 |
| Thr | Leu | Glu | Leu | Arg<br>485 | Arg | Val | Thr  | Ala | Glu<br>490 | Glu | Ala | Gly | Leu | Tyr<br>495 |
| Thr | Cys | Val | Ala | Gln<br>500 | Asn | Leu | Val  | Gly | Ala<br>505 | Asp | Thr | Lys | Thr | Val<br>510 |
| Ser | Val | Val | Val | Gly<br>515 | Arg | Ala | Leu  | Leu | Gln<br>520 | Pro | Gly | Arg | Asp | Glu<br>525 |
| Gly | Gln | Gly | Leu | Glu<br>530 | Leu | Arg | Val  | Gln | Glu<br>535 | Thr | His | Pro | Tyr | His<br>540 |
| Ile | Leu | Leu | Ser | Trp<br>545 | Val | Thr | Pro  | Pro | Asn<br>550 | Thr | Val | Ser | Thr | Asn<br>555 |
| Leu | Thr | Trp | Ser | Ser<br>560 | Ala | Ser | Ser  | Leu | Arg<br>565 | Gly | Gln | Gly | Ala | Thr<br>570 |
| Ala | Leu | Ala | Arg | Leu<br>575 | Pro | Arg | Gly  | Thr | His<br>580 | Ser | Tyr | Asn | Ile | Thr<br>585 |
| Arg | Leu | Leu | Gln | Ala<br>590 | Thr | Glu | Tyr  | Trp | Ala<br>595 | Сув | Leu | Gln | Val | Ala<br>600 |
| Phe | Ala | Asp | Ala | His<br>605 | Thr | Gln | Leu  | Ala | Cys<br>610 | Val | Trp | Ala | Arg | Thr<br>615 |
| Lys | Glu | Ala | Thr | Ser<br>620 | Сув | His | Arg  | Ala | Leu<br>625 | Gly | Asp | Arg | Pro | Gly<br>630 |
| Leu | Ile | Ala | Ile | Leu<br>635 | Ala | Leu | Ala  | Val | Leu<br>640 | Leu | Leu | Ala | Ala | Gly<br>645 |
| Leu | Ala | Ala | His | Leu<br>650 | Gly | Thr | Gly. | Gln | Pro<br>655 | Arg | Lys | Gly | Val | Gly<br>660 |
| Gly | Arg | Arg | Pro | Leu<br>665 | Pro | Pro | Ala  | Trp | Ala<br>670 | Phe | Trp | Gly | Trp | Ser        |

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Cys Ala His Pro Leu Ala Thr Leu Phe Lys Ile Leu Ala Ser Phe
50 55 60

Tyr Ile Ser Leu Val Ile Phe Tyr Gly Leu Ile Cys Met Tyr Thr
65 70 75

Leu Trp Trp Met Leu Arg Arg Ser Leu Lys Lys Tyr Ser Phe Glu 80 85 90

Ser Ile Arg Glu Glu Ser Ser Tyr Ser Asp Ile Pro Asp Val Lys





|     |     |     |     | 95          |     |     |     |     | 100        |     |     |     |          | 105        |
|-----|-----|-----|-----|-------------|-----|-----|-----|-----|------------|-----|-----|-----|----------|------------|
| Asn | Asp | Phe | Ala | Phe<br>110  | Met | Leu | His | Leu | Ile<br>115 | Asp | Gln | Tyr | Asp      | Pro<br>120 |
| Leu | Tyr | Ser | Lys | Arg<br>125  | Phe | Ala | Val | Phe | Leu<br>130 | Ser | Glu | Val | Ser      | Glu<br>135 |
| Asn | Lys | Leu | Arg | Gln<br>140  | Leu | Asn | Leu | Asn | Asn<br>145 | Glu | Trp | Thr | Leu      | Asp<br>150 |
| Lys | Leu | Arg | Gln | Arg<br>155  | Leu | Thr | Lys | Asn | Ala<br>160 | Gln | Asp | Lys | Leu      | Glu<br>165 |
| Leu | His | Leu | Phe | Met<br>170  | Leu | Ser | Gly | Ile | Pro<br>175 | Asp | Thr | Val | Phe      | Asp<br>180 |
| Leu | Val | Glu | Leu | Glu<br>185  | Val | Leu | Lys | Leu | Glu<br>190 | Leu | Ile | Pro | Asp      | Val<br>195 |
| Thr | Ile | Pro | Pro | Ser<br>200  | Ile | Ala | Gln | Leu | Thr<br>205 | Gly | Leu | Lys | Glu      | Leu<br>210 |
| Trp | Leu | Tyr | His | Thir<br>215 | Ala | Ala | Lys | Ile | Glu<br>220 | Ala | Pro | Ala | Leu      | Ala<br>225 |
| Phe | Leu | Arg | Glu | Asn<br>230  | Leu | Arg | Ala | Leu | His<br>235 | Ile | Lys | Phe | Thr      | Asp<br>240 |
| Ile | Lys | Glu | Ile | Pro<br>245  | Leu | Trp | Ile | Tyr | Ser<br>250 | Leu | Lys | Thr | Leu      | Glu<br>255 |
| Glu | Leu | His | Leu | Thr<br>260  | Gly | Asn | Leu | Ser | Ala<br>265 | Glu | Asn | Asn | Arg      | Tyr<br>270 |
| Ile | Val | Ile | Asp | Gly<br>275  | Leu | Arg | Glu | Leu | Lys<br>280 | Arg | Leu | Lys | Val      | Leu<br>285 |
| Arg | Leu | Lys | Ser |             | Leu | Ser | Lys | Leu | Pro<br>295 | Gln | Val | Val | Thr      | Asp<br>300 |
| Val | Glv | Val | His | 290<br>Leu  | Gln | Lvs | Leu | Ser |            | Asn | Asn | Glu | Gly      |            |
|     | 1   |     |     | 305         |     | -1- |     |     | 310        |     |     |     | <b>4</b> | 315        |
| Lys | Leu | Ile | Val | Leu<br>320  | Asn | Ser | Leu | Lys | Lys<br>325 | Met | Ala | Asn | Leu      | Thr<br>330 |
| Glu | Leu | Glu | Leu | 11e<br>335  | Arg | Cys | Asp | Leu | Glu<br>340 | Arg | Ile | Pro | His      | Ser<br>345 |
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<sup>&</sup>lt;210> 255

<sup>&</sup>lt;211> 452

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

<sup>&</sup>lt;400> 255

Met Glu Leu Ala Leu Arg Arg Ser Pro Val Pro Arg Trp Leu Leu

1 5 10 15

Leu Leu Pro Leu Leu Gly Leu Asn Ala Gly Ala Val Ile Asp

Trp Pro Thr Glu Glu Gly Lys Glu Val Trp Asp Tyr Val Thr Val
35 40 45

| Arg    | Lys | Asp | Ala                | Tyr<br>50  | Met | Phe | Trp | Trp | Leu<br>55  | Tyr | Tyr | Ala | Thr | Asn<br>60          |
|--------|-----|-----|--------------------|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|--------------------|
| Ser    | Cys | Lys | Asn                | Phe<br>65  | Ser | Glu | Leu | Pro | Leu<br>70  | Val | Met | Trp | Leu | Gln<br>75          |
| Gly    | Gly | Pro | Gly                | Gly<br>80  | Ser | Ser | Thr | Gly | Phe<br>85  | Gly | Asn | Phe | Glu | Glu<br>90          |
| Ile    | Gly | Pro | Leu                | Asp<br>95  | Ser | Asp | Leu | Lys | Pro<br>100 | Arg | Lys | Thr | Thr | Trp<br>105         |
| Leu    | Gln | Ala | Ala                | Ser<br>110 | Leu | Leu | Phe | Val | Asp<br>115 | Asn | Pro | Val | Gly | Thr<br>120         |
| Gly    | Phe | Ser | Tyr                | Val<br>125 | Asn | Gly | Ser | Gly | Ala<br>130 | Tyr | Ala | Lys | Asp | Leu<br><b>1</b> 35 |
| Ala    | Met | Val | Ala                | Ser<br>140 | Asp | Met | Met | Val | Leu<br>145 | Leu | Lys | Thr | Phe | Phe<br>150         |
| Ser    | Cys | His | Lys                | Glu<br>155 | Phe | Gln | Thr | Val | Pro<br>160 | Phe | Tyr | Ile | Phe | Ser<br>165         |
| Glu    | Ser | Tyr | Gly                | Gly<br>170 | Lys | Met | Ala | Ala | Gly<br>175 | Ile | Gly | Leu | Glu | Leu<br>180         |
| Tyr    | Lys | Ala | Ile                | Gln<br>185 | Arg | Gly | Thr | Ile | Lys<br>190 | Cys | Asn | Phe | Ala | Gly<br>195         |
| Val    | Ala | Leu | Gly                | Asp<br>200 | Ser | Trp | Ile | Ser | Pro<br>205 | Val | Asp | Ser | Val | Leu<br>210         |
| Ser    | Trp | Gly | Pro                | Tyr<br>215 | Leu | Tyr | Ser | Met | Ser<br>220 | Leu | Leu | Glu | Asp | Lys<br>225         |
| Gly    | Leu | Ala | Glu                | Val<br>230 | Ser | Lys | Val | Ala | Glu<br>235 | Gln | Val | Leu | Asn | Ala<br>240         |
| Val    | Asn | Lys | Gly                | Leu<br>245 | Tyr | Arg | Glu | Ala | Thr<br>250 | Glu | Leu | Trp | Gly | Lys<br>255         |
| Ala    | Glu | Met | Ile                | Ile<br>260 | Glu | Gln | Asn | Thr | Asp<br>265 | Gly | Val | Asn | Phe | Tyr<br>270         |
| Asn    | Ile | Leu | Thr                | Lys<br>275 | Ser | Thr | Pro | Thr | Ser<br>280 | Thr | Met | Glu | Ser | Ser<br>285         |
| Leu    | Glu | Phe | Thr                | Gln<br>290 | Ser | His | Leu | Val | Cys<br>295 | Leu | Cys | Gln | Arg | His<br>300         |
| 17 a 1 | Ara | Hic | T. <del>2</del> 11 | Gln        | Ara | Asn | Δla | Len | Ser        | Gln | Leu | Met | Asn | Gly                |

|   | 305               |        |       |      | 310        |      |     |      |     | 315        |
|---|-------------------|--------|-------|------|------------|------|-----|------|-----|------------|
| Pro Ile Arg Ly  | s Lys Leu<br>320  | Lys    | Ile   | Ile  | Pro<br>325 | Glu  | Asp | Gln  | Ser | Trp<br>330 |
| Gly Gly Gln Al  | a Thr Asn<br>335  | Val    | Phe   | Val  | Asn<br>340 | Met  | Glu | Glu  | Asp | Phe<br>345 |
| Met Lys Pro Va  | l Ile Ser<br>350  | Ile    | Val   | Asp  | Glu<br>355 | Leu  | Leu | Glu  | Ala | Gly<br>360 |
| Ile Asn Val Th  | r Val Tyr<br>365  | Asn    | Gly   | Gln  | Leu<br>370 | Asp  | Leu | Ile  | Val | Asp<br>375 |
| Thr Met Gly Gl  | n Glu Ala<br>380  | Trp    | Val   | Arg  | Lys<br>385 | Leu  | Lys | Trp  | Pro | Glu<br>390 |
| Leu Pro Lys Ph  | e Ser Gln<br>395  | Leu    | Lys   | Trp  | Lys<br>400 | Ala  | Leu | Tyr  | Ser | Asp<br>405 |
| Pro Lys Ser Le  | u Glu Thr<br>410  | Ser    | Ala   | Phe  | Val<br>415 | Lys  | Ser | Tyr  | Lys | Asn<br>420 |
| Leu Ala Phe Ty  | r Trp Ile<br>425  | Leu    | Lys   | Ala  | Gly<br>430 | His  | Met | Val  | Pro | Ser<br>435 |
| Asp Gln Gly As  | p Met Ala<br>440  | Leu    | Lys   | Met  | Met<br>445 | Arg  | Leu | Val  | Thr | Gln<br>450 |
| Gln Glu   |                   |        |       |      |            |      |     |      |     |            |
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| ccgttatcag gac  | catgcgg c         | cgacç  | gggto | c at | cacg       | tcgc | gca | tcgt | ggg | 150        |
| tggagaggac gcc  | gaactcg g         | gcgtt  | ggc   | g gt | ggca       | gggg | agc | ctgc | gcc | 200        |
| tgtgggattc cca  | .cgtatgc <u>c</u> | gagto  | gagc  | c tg | ctca       | gcca | ccg | ctgg | gca | 250        |
| ctcacggcgg cgc  | actgctt t         | gaaa   | ccta  | t ag | tgac       | ctta | gtg | atcc | ctc | 300        |
| cgggtggatg gto  | cagtttg g         | gccago | ctga  | c tt | ccat       | gcca | tcc | ttct | gga | 350        |

gcctgcaggc ctactacacc cgttacttcg tatcgaatat ctatctgagc 400

<400> 257

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1 5 10 15

Ala Gly Leu Arg Lys Pro Glu Ser Gln Glu Ala Ala Pro Leu Ser

20 25 30

Gly Pro Cys Gly Arg Arg Val Ile Thr Ser Arg Ile Val Gly Gly
35 40 45

Glu Asp Ala Glu Leu Gly Arg Trp Pro Trp Gln Gly Ser Leu Arg
50 55 60

Leu Trp Asp Ser His Val Cys Gly Val Ser Leu Leu Ser His Arg

Trp Ala Leu Thr Ala Ala His Cys Phe Glu Thr Tyr Ser Asp Leu
80 85 90

<sup>&</sup>lt;210> 257

<sup>&</sup>lt;211> 314

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

| Ser   | Asp  | Pro | Ser | Gly<br>95  | Trp | Met | Val | Gln | Phe<br>100 | Gly | Gln | Leu | Thr | Ser<br>105 |
|-------|------|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Met   | Pro  | Ser | Phe | Trp<br>110 | Ser | Leu | Gln | Ala | Tyr<br>115 | Tyr | Thr | Arg | Tyr | Phe<br>120 |
| Val   | Ser  | Asn | Ile | Tyr<br>125 | Leu | Ser | Pro | Arg | Tyr<br>130 | Leu | Gly | Asn | Ser | Pro<br>135 |
| Tyr   | Asp  | Ile | Ala | Leu<br>140 | Val | Lys | Leu | Ser | Ala<br>145 | Pro | Val | Thr | Tyr | Thr<br>150 |
| Lys   | His  | Ile | Gln | Pro<br>155 | Ile | Cys | Leu | Gln | Ala<br>160 | Ser | Thr | Phe | Glu | Phe<br>165 |
| Glu   | Asn  | Arg | Thr | Asp<br>170 | Cys | Trp | Val | Thr | Gly<br>175 | Trp | Gly | Tyr | Ile | Lys<br>180 |
| Glu   | Asp  | Glu | Ala | Leu<br>185 | Pro | Ser | Pro | His | Thr<br>190 | Leu | Gln | Glu | Val | Gln<br>195 |
| Val   | Ala  | Ile | Ile | Asn<br>200 | Asn | Ser | Met | Cys | Asn<br>205 | His | Leu | Phe | Leu | Lys<br>210 |
| Tyr   | Ser  | Phe | Arg | Lys<br>215 | Asp | Ile | Phe | Gly | Asp<br>220 | Met | Val | Cys | Ala | Gly<br>225 |
| Asn   | Ala  | Gln | Gly | Gly<br>230 | Lys | Asp | Ala | Cys | Phe<br>235 | Gly | Asp | Ser | Gly | Gly<br>240 |
| Pro   | Leu  | Ala | Cys | Asn<br>245 | Lys | Asn | Gly | Leu | Trp<br>250 | Tyr | Gln | Ile | Gly | Val<br>255 |
| Val   | Ser  | Trp | Gly | Val<br>260 | Gly | Cys | Gly | Arg | Pro<br>265 | Asn | Arg | Pro | Gly | Val<br>270 |
| Tyr   | Thr  | Asn | Ile | Ser<br>275 | His | His | Phe | Glu | Trp<br>280 | Ile | Gln | Lys | Leu | Met<br>285 |
| Ala   | Gln  | Ser | Gly | Met<br>290 | Ser | Gln | Pro | Asp | Pro<br>295 | Ser | Trp | Pro | Leu | Leu<br>300 |
| Phe   | Phe  | Pro | Leu | Leu<br>305 | Trp | Ala | Leu | Pro | Leu<br>310 | Leu | Gly | Pro | Val |            |
| <210: | > 25 | 8   |     |            |     |     |     |     |            | •   |     |     |     |            |

<sup>&</sup>lt;210> 258

<sup>&</sup>lt;211> 2427

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo Sapien

<sup>&</sup>lt;400> 258

cccacgcgtc cgcggacgcg tgggaagggc agaatgggac tccaagcctg 50

cctcctaggg ctctttgccc tcatcctctc tggcaaatgc agttacagcc 100 eggageeega eeageggagg aegetgeeee eaggetgggt gteeetggge 150 cgtgcggacc ctgaggaaga gctgagtctc acctttgccc tgagacagca 200 gaatgtggaa agactctcgg agctggtgca ggctgtgtcg gatcccagct 250 ctcctcaata cggaaaatac ctgaccctag agaatgtggc tgatctggtg 300 aggecatece caetgaceet ceacaeggtg caaaaatgge tettggeage 350 cggagcccag aagtgccatt ctgtgatcac acaggacttt ctgacttgct 400 ggctgagcat ccgacaagca gagctgctgc tccctggggc tgagtttcat 450 cactatgtgg gaggacctac ggaaacccat gttgtaaggt ccccacatcc 500 ctaccagett ccacaggeet tggeeceeca tgtggaettt gtggggggae 550 tgcaccgttt tcccccaaca tcatccctga ggcaacgtcc tgagccgcag 600 gtgacaggga ctgtaggcct gcatctgggg gtaaccccct ctgtgatccg 650 taagcgatac aacttgacct cacaagacgt gggctctggc accagcaata 700 acagccaagc ctgtgcccag ttcctggagc agtatttcca tgactcagac 750 ctggctcagt tcatgcgcct cttcggtggc aactttgcac atcaggcatc 800 ccagtctaga tgtgcagtac ctgatgagtg ctggtgccaa catctccacc 900 tgggtctaca gtagccctgg ccggcatgag ggacaggagc ccttcctgca 950 qtqqctcatq ctgctcagta atgagtcagc cctgccacat gtgcatactg 1000 tgagctatgg agatgatgag gactccctca gcagcgccta catccagcgg 1050 gtcaacactg agctcatgaa ggctgccgct cggggtctca ccctgctctt 1100 cgcctcaggt gacagtgggg ccgggtgttg gtctgtctct ggaagacacc 1150 agttccgccc taccttccct gcctccagcc cctatgtcac cacagtggga 1200 ggcacatcct tccaggaacc tttcctcatc acaaatgaaa ttgttgacta 1250 tatcagtggt ggtggcttca gcaatgtgtt cccacggcct tcataccagg 1300 aggaagetgt aacgaagtte etgageteta gececeaeet gecaceatee 1350 agttacttca atgccagtgg ccgtgcctac ccagatgtgg ctgcactttc 1400

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Ser Gly Lys Cys Ser Tyr Ser Pro Glu Pro Asp Gln Arg Arg Thr

<sup>&</sup>lt;210> 259

<sup>&</sup>lt;211> 556

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

<sup>&</sup>lt;400> 259

Met Gly Leu Gln Ala Cys Leu Leu Gly Leu Phe Ala Leu Ile Leu

1 5 10 15

|     |     |     |     | 20         |     |     |     |     | 25         |     |     |     |     | 30         |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Leu | Pro | Pro | Gly | Trp<br>35  | Val | Ser | Leu | Gly | Arg<br>40  | Ala | Asp | Pro | Glu | Glu<br>45  |
| Glu | Leu | Ser | Leu | Thr<br>50  | Phe | Ala | Leu | Arg | Gln<br>55  | Gln | Asn | Val | Glu | Arg<br>60  |
| Leu | Ser | Glu | Leu | Val<br>65  | Gln | Ala | Val | Ser | Asp<br>70  | Pro | Ser | Ser | Pro | Gln<br>75  |
| Tyr | Gly | Lys | Tyr | Leu<br>80  | Thr | Leu | Glu | Asn | Val<br>85  | Ala | Asp | Leu | Val | Arg<br>90  |
| Pro | Ser | Pro | Leu | Thr<br>95  | Leu | His | Thr | Val | Gln<br>100 | Lys | Trp | Leu | Leu | Ala<br>105 |
| Ala | Gly | Ala | Gln | Lys<br>110 | Суз | His | Ser | Val | Ile<br>115 | Thr | Gln | Asp | Phe | Leu<br>120 |
| Thr | Cys | Trp | Leu | Ser<br>125 | Ile | Arg | Gln | Ala | Glu<br>130 | Leu | Leu | Leu | Pro | Gly<br>135 |
| Ala | Glu | Phe | His | His<br>140 | Tyr | Val | Gly | Gly | Pro<br>145 | Thr | Glu | Thr | His | Val<br>150 |
| Val | Arg | Ser | Pro | His<br>155 | Pro | Tyr | Gln | Leu | Pro<br>160 | Gln | Ala | Leu | Ala | Pro<br>165 |
| His | Val | Asp | Phe | Val<br>170 | Gly | Gly | Leu | His | Arg<br>175 | Phe | Pro | Pro | Thr | Ser<br>180 |
| Ser | Leu | Arg | Gln | Arg<br>185 | Pro | Glu | Pro | Gln | Val<br>190 | Thr | Gly | Thr | Val | Gly<br>195 |
| Leu | His | Leu | Gly | Val<br>200 | Thr | Pro | Ser | Val | Ile<br>205 | Arg | Lys | Arg | Tyr | Asn<br>210 |
| Leu | Thr | Ser |     | Asp<br>215 |     | Gly | Ser | Gly | Thr<br>220 |     | Asn | Asn | Ser | Gln<br>225 |
| Ala | Cys | Ala | Gln | Phe<br>230 |     | Glu | Gln | Tyr | Phe<br>235 |     | Asp | Ser | Asp | Leu<br>240 |
| Ala | Gln | Phe | Met | Arg<br>245 |     | Phe | Gly | Gly | Asn<br>250 |     | Ala | His | Gln | Ala<br>255 |
| Ser | Val | Ala | Arg | Val<br>260 |     | Gly | Gln | Gln | Gly<br>265 |     | Gly | Arg | Ala | Gly<br>270 |
| Ile | Glu | Ala | Ser | Leu<br>275 |     | Val | Gln | Tyr | Leu<br>280 |     | Ser | Ala | Gly | Ala<br>285 |

| Asn | Ile | Ser | Thr | Trp<br>290 | Val | туr | Ser | Ser | Pro<br>295 | Gly | Arg | His | Glu | Gly<br>300 |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Gln | Glu | Pro | Phe | Leu<br>305 | Gln | Trp | Leu | Met | Leu<br>310 | Leu | Ser | Asn | Glu | Ser<br>315 |
| Ala | Leu | Pro | His | Val<br>320 | His | Thr | Val | Ser | Tyr<br>325 | Gly | Asp | Asp | Glu | Asp<br>330 |
| Ser | Leu | Ser | Ser | Ala<br>335 | Tyr | Ile | Gln | Arg | Val<br>340 | Asn | Thr | Glu | Leu | Met<br>345 |
| Lys | Ala | Ala | Ala | Arg<br>350 | Gly | Leu | Thr | Leu | Leu<br>355 | Phe | Ala | Ser | Gly | Asp<br>360 |
| Ser | Gly | Ala | Gly | Cys<br>365 | Trp | Ser | Val | Ser | Gly<br>370 | Arg | His | Gln | Phe | Arg<br>375 |
| Pro | Thr | Phe | Pro | Ala<br>380 | Ser | Ser | Pro | Tyr | Val<br>385 | Thr | Thr | Val | Gly | Gly<br>390 |
| Thr | Ser | Phe | Gln | Glu<br>395 | Pro | Phe | Leu | Ile | Thr<br>400 | Asn | Glu | Ile | Val | Asp<br>405 |
| Tyr | Ile | Ser | Gly | Gly<br>410 | Gly | Phe | Ser | Asn | Val<br>415 | Phe | Pro | Arg | Pro | Ser<br>420 |
| Tyr | Gln | Glu | Glu | Ala<br>425 | Val | Thr | Lys | Phe | Leu<br>430 | Ser | Ser | Ser | Pro | His<br>435 |
| Leu | Pro | Pro | Ser | Ser<br>440 | Tyr | Phe | Asn | Ala | Ser<br>445 | Gly | Arg | Ala | Tyr | Pro<br>450 |
| Asp | Val | Ala | Ala | Leu<br>455 | Ser | Asp | Gly | Tyr | Trp<br>460 | Val | Val | Ser | Asn | Arg<br>465 |
| Val | Pro | Ile | Pro | Trp<br>470 | Val | Ser | Gly | Thr | Ser<br>475 | Ala | Ser | Thr | Pro | Val<br>480 |
| Phe | Gly | Gly | Ile | Leu<br>485 | Ser | Leu | Ile | Asn | Glu<br>490 | His | Arg | Ile | Leu | Ser<br>495 |
| Gly | Arg | Pro | Pro | Leu<br>500 | Gly | Phe | Leu | Asn | Pro<br>505 | Arg | Leu | Tyr | Gln | Gln<br>510 |
| His | Gly | Ala | Gly | Leu<br>515 | Phe | Asp | Val | Thr | Arg<br>520 | Gly | Cys | His | Glu | Ser<br>525 |
| Cys | Leu | Asp | Glu | Glu<br>530 | Val | Glu | Gly | Gln | Gly<br>535 | Phe | Cys | Ser | Gly | Pro<br>540 |
| Gly | Trp | Asp | Pro | Val<br>545 | Thr | Gly | Trp | Gly | Thr<br>550 | Pro | Thr | Ser | Gln | Leu<br>555 |

Cys

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<213> Homo Sapien

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<211> 383

<212> PRT

<213> Homo Sapien

<400> 261

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Cys Ala Val Gly Gln Val Ser Pro Tyr Ser Ala Pro Trp Lys Pro  $20 \hspace{1cm} 25 \hspace{1cm} 30 \hspace{1cm}$ 

Thr Trp Pro Ala Tyr Arg Leu Pro Val Val Leu Pro Gln Ser Thr 35 40 45

Leu Asn Leu Ala Lys Pro Asp Phe Gly Ala Glu Ala Lys Leu Glu
50 55 60

Val Ser Ser Cys Gly Pro Gln Cys His Lys Gly Thr Pro Leu 65 70 75

Pro Thr Tyr Glu Glu Ala Lys Gln Tyr Leu Ser Tyr Glu Thr Leu 80 85 90

Tyr Ala Asn Gly Ser Arg Thr Glu Thr Gln Val Gly Ile Tyr Ile

95 100 105

Leu Ser Ser Ser Gly Asp Gly Ala Gln His Arg Asp Ser Gly Ser 110 115 120

| Ser | Gly | Lys | Ser | Arg<br>125 | Arg | Lys | Arg | Gln | Ile<br>130 | Tyr | Gly | Tyr | Asp   | Ser<br>135 |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-------|------------|
| Arg | Phe | Ser | Ile | Phe<br>140 | Gly | Lys | Asp | Phe | Leu<br>145 | Leu | Asn | Tyr | Pro   | Phe<br>150 |
| Ser | Thr | Ser | Val | Lys<br>155 | Leu | Ser | Thr | Gly | Cys<br>160 | Thr | Gly | Thr | Leu   | Val<br>165 |
| Ala | Glu | Lys | His | Val<br>170 | Leu | Thr | Ala | Ala | His<br>175 | Cys | Ile | His | Asp   | Gly<br>180 |
| Lys | Thr | Tyr | Val | Lys<br>185 | Gly | Thr | Gln | Lys | Leu<br>190 | Arg | Val | Gly | Phe   | Leu<br>195 |
| Lys | Pro | Lys | Phe | Lys<br>200 | Asp | Gly | Gly | Arg | Gly<br>205 | Ala | Asn | Asp | Ser   | Thr<br>210 |
| Ser | Ala | Met | Pro | Glu<br>215 | Gln | Met | Lys | Phe | Gln<br>220 | Trp | Ile | Arg | Val   | Lys<br>225 |
| Arg | Thr | His | Val | Pro<br>230 | Lys | Gly | Trp | Ile | Lys<br>235 | Gly | Asn | Ala | Asn   | Asp<br>240 |
| Ile | Gly | Met | Asp | Tyr<br>245 | Asp | Tyr | Ala | Leu | Leu<br>250 | Glu | Leu | Lys | Lys   | Pro<br>255 |
| His | Lys | Arg | Lys | Phe<br>260 | Met | Lys | Ile | Gly | Val<br>265 | Ser | Pro | Pro | Ala   | Lys<br>270 |
| Gln | Leu | Pro | Gly | Gly<br>275 | Arg | Ile | His | Phe | Ser<br>280 | Gly | Tyr | Asp | Asn   | Asp<br>285 |
| Arg | Pro | Gly | Asn | Leu<br>290 | Val | Tyr | Arg | Phe | Cys<br>295 | Asp | Val | Lys | Asp   | Glu<br>300 |
| Thr | Tyr | Asp | Leu | Leu<br>305 | Tyr | Gln | Gln | Cys | Asp<br>310 | Ala | Gln | Pro | Gly   | Ala<br>315 |
| Ser | Gly | Ser | Gly | Val<br>320 | Tyr | Val | Arg | Met | Trp<br>325 | Lys | Arg | Gln | Gln   | Gln<br>330 |
| Lys | Trp | Glu | Arg | Lys<br>335 | Ile | Ile | Gly | Ile | Phe<br>340 |     | Gly | His | Gln   | Trp<br>345 |
| Val | Asp | Met | Asn | Gly<br>350 |     | Pro | Gln | Asp | Phe<br>355 |     | Val | Ala | . Val | Arg<br>360 |
| Ile | Thr | Pro | Leu | Lys<br>365 |     | Ala | Gln | Ile | Cys<br>370 |     | Trp | Ile | . Lys | Gly<br>375 |
| Asn | туг | Leu | Asp | Cys<br>380 |     | Glu | Gly | •   |            |     |     |     |       |            |

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aggaaaccc ctccccgacc cgcccgacg cctcaggcc ccctccaagg 1200 catcaggcc cgcccaacgg cctcatgtcc ccgccccac gacttccggc 1250 cccgccccc ggccccagcg cttttgtgta tataaatgtt aatgatttt 1300 ataggtattt gtaaccctgc ccacatatct tatttattcc tccaatttca 1350 ataaattatt tattctccaa aaaaaaaa 1378

<210> 263

<211> 317

<212> PRT

<213> Homo Sapien

<400> 263

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170

175

180

## 

| Gly          | Ser           | Ile            | Gln   | Asp<br>185 | Gly   | Val    | Pro   | Leu     | Pro<br>190 | His | Pro | Gln | Thr | Leu<br>195 |
|--------------|---------------|----------------|-------|------------|-------|--------|-------|---------|------------|-----|-----|-----|-----|------------|
| Gln          | Lys           | Leu            | Lys   | Val<br>200 | Pro   | Ile    | Ile   | Asp     | Ser<br>205 | Glu | Val | Cys | Ser | His<br>210 |
| Leu          | Tyr           | Trp            | Arg   | Gly<br>215 | Ala   | Gly    | Gln   | Gly     | Pro<br>220 | Ile | Thr | Glu | Asp | Met<br>225 |
| Leu          | Cys           | Ala            | Gly   | Tyr<br>230 | Leu   | Glu    | Gly   | Glu     | Arg<br>235 | Asp | Ala | Cys | Leu | Gly<br>240 |
| Asp          | Ser           | Gly            | Gly   | Pro<br>245 | Leu   | Met    | Cys   | Gln     | Val<br>250 | Asp | Gly | Ala | Trp | Leu<br>255 |
| Leu          | Ala           | Gly            | Ile   | Ile<br>260 | Ser   | Trp    | Gly   | Glu     | Gly<br>265 | Cys | Ala | Glu | Arg | Asn<br>270 |
| Arg          | Pro           | Gly            | Val   | Tyr<br>275 | Ile   | Ser    | Leu   | Ser     | Ala<br>280 | His | Arg | Ser | Trp | Val<br>285 |
| Glu          | Lys           | Ile            | Val   | Gln<br>290 | Gly   | Val    | Gln   | Leu     | Arg<br>295 | Gly | Arg | Ala | Gln | Gly<br>300 |
| Gly          | Gly           | Ala            | Leu   | Arg<br>305 | Ala   | Pro    | Ser   | Gln     | Gly<br>310 | Ser | Gly | Ala | Ala | Ala<br>315 |
| Arg          | Ser           |                |       |            |       |        |       |         |            |     |     |     |     |            |
| 211:         | DNA           | Ą              | cial  | Sequ       | ience | e      |       |         |            |     |     |     |     |            |
| 220:         |               | - <del>-</del> |       | \] d ~~    |       | loot i | ido T | ) wah a |            |     |     |     |     |            |
| (223)        | > Syr         | ıcnet          | ite ( | )11gc      | nucl  | Leot   | ide i | Probe   | 3          |     |     |     |     |            |
|              | > 264<br>gcaa |                | atgco | etaca      | at gt | tc 2   | 24    |         |            |     |     |     |     |            |
| 211<br>212   | DNA           | Ā              | cial  | Sequ       | ience | e      |       |         |            |     |     |     |     |            |
| :220<br>:223 |               | nthet          | cic ( | )ligo      | nucl  | leoti  | ide I | Probe   | <u> </u>   |     |     |     |     |            |
|              | 265<br>Jaggt  |                | ctaaç | gttg       | g 19  |        |       |         |            |     |     |     |     |            |
| 210          | 266<br>24     | 5              |       |            |       |        |       |         |            |     |     |     |     |            |

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<400> 266
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<400> 267
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 cagccctgcc acatgtgc 18
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Glu Glu Lys Arg Leu Met Val Glu Leu His Asn Leu Tyr Arg Ala 35 40 45

Gln Val Ser Pro Thr Ala Ser Asp Met Leu His Met Arg Trp Asp
50 55 60

Glu Glu Leu Ala Ala Phe Ala Lys Ala Tyr Ala Arg Gln Cys Val 65 70 75

Trp Gly His Asn Lys Glu Arg Gly Arg Arg Gly Glu Asn Leu Phe

Ala Ile Thr Asp Glu Gly Met Asp Val Pro Leu Ala Met Glu Glu 95 100 105

Trp His His Glu Arg Glu His Tyr Asn Leu Ser Ala Ala Thr Cys 110 115 120

Ser Pro Gly Gln Met Cys Gly His Tyr Thr Gln Val Val Trp Ala

| Lys | Thr | Glu | Arg | Ile<br>140 | Gly | Cys | Gly | Ser | His<br>145 | Phe | Cys | Glu | Lys | Leu<br>150 |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Gln | Gly | Val | Glu | Glu<br>155 | Thr | Asn | Ile | Glu | Leu<br>160 | Leu | Val | Cys | Asn | Tyr<br>165 |
| Glu | Pro | Pro | Gly | Asn<br>170 | Val | Lys | Gly | Lys | Arg<br>175 | Pro | Tyr | Gln | Glu | Gly<br>180 |
| Thr | Pro | Cys | Ser | Gln<br>185 | Cys | Pro | Ser | Gly | Tyr<br>190 | His | Cys | Lys | Asn | Ser<br>195 |
| Leu | Cys | Glu | Pro | Ile<br>200 | Gly | Ser | Pro | Glu | Asp<br>205 | Ala | Gln | Asp | Leu | Pro<br>210 |
| Tyr | Leu | Val | Thr | Glu<br>215 | Ala | Pro | Ser | Phe | Arg<br>220 | Ala | Thr | Glu | Ala | Ser<br>225 |
| Asp | Ser | Arg | Lys | Met<br>230 | Gly | Thr | Pro | Ser | Ser<br>235 | Leu | Ala | Thr | Gly | Ile<br>240 |
| Pro | Ala | Phe | Leu | Val<br>245 | Thr | Glu | Val | Ser | Gly<br>250 | Ser | Leu | Ala | Thr | Lys<br>255 |
| Ala | Leu | Pro | Ala | Val<br>260 | Glu | Thr | Gln | Ala | Pro<br>265 | Thr | Ser | Leu | Ala | Thr<br>270 |
| Lys | Asp | Pro | Pro | Ser<br>275 | Met | Ala | Thr | Glu | Ala<br>280 | Pro | Pro | Cys | Val | Thr<br>285 |
| Thr | Glu | Val | Pro | Ser<br>290 | Ile | Leu | Ala | Ala | His<br>295 | Ser | Leu | Pro | Ser | Leu<br>300 |
| Asp | Glu | Glu | Pro | Val<br>305 | Thr | Phe | Pro | Lys | Ser<br>310 | Thr | His | Val | Pro | Ile<br>315 |
| Pro | Lys | Ser | Ala | Asp<br>320 | Lys | Val | Thr | Asp | Lys<br>325 | Thr | Lys | Val | Pro | Ser<br>330 |
| Arg | Ser | Pro | Glu | Asn<br>335 | Ser | Leu | Asp | Pro | Lys<br>340 | Met | Ser | Leu | Thr | Gly<br>345 |
| Ala | Arg | Glu | Leu | Leu<br>350 | Pro | His | Ala | Gln | Glu<br>355 | Glu | Ala | Glu | Ala | Glu<br>360 |
| Ala | Glu | Leu | Pro | Pro<br>365 | Ser | Ser | Glu | Val | Leu<br>370 | Ala | Ser | Val | Phe | Pro<br>375 |
| Ala | Gln | Asp | Lys | Pro<br>380 | Gly | Glu | Leu | Gln | Ala<br>385 | Thr | Leu | Asp | His | Thr<br>390 |
| Gly | His | Thr | Ser | Ser<br>395 | Lys | Ser | Leu | Pro | Asn<br>400 | Phe | Pro | Asn | Thr | Ser<br>405 |

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Gly Leu Asn Ser Gly Pro Gly His Val Trp Gly Pro Leu Leu Gly
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<400> 290

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Ile Ser Arg Pro Asp Leu Ser His Asn Arg Leu Ser Phe Ile Lys
20 25 30

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|--------|--------|-------|-----|------------|-----|-----|-----|-----|------------|-----|-----|-------|-----|------------|
| Asn    | Asn    | Asn   | Glu | Leu<br>50  | Glu | Thr | Ile | Pro | Asn<br>55  | Leu | Gly | Pro   | Val | Ser<br>60  |
| Ala    | Asn    | Ile   | Thr | Leu<br>65  | Leu | Ser | Leu | Ala | Gly<br>70  | Asn | Arg | Ile   | Val | Glu<br>75  |
| Ile    | Leu    | Pro   | Glu | His<br>80  | Leu | Lys | Glu | Phe | Gln<br>85  | Ser | Leu | Glu   | Thr | Leu<br>90  |
| Asp    | Leu    | Ser   | Ser | Asn<br>95  | Asn | Ile | Ser | Glu | Leu<br>100 | Gln | Thr | Ala   | Phe | Pro<br>105 |
| Ala    | Leu    | Gln   | Leu | Lys<br>110 | Tyr | Leu | Tyr | Leu | Asn<br>115 | Ser | Asn | Arg   | Val | Thr<br>120 |
| Ser    | Met    | Glu   | Pro | Gly<br>125 | Tyr | Phe | Asp | Asn | Leu<br>130 | Ala | Asn | Thr   | Leu | Leu<br>135 |
| Val    | Leu    | Lys   | Leu | Asn<br>140 | Arg | Asn | Arg | Ile | Ser<br>145 | Ala | Ile | Pro   | Pro | Lys<br>150 |
| Met    | Phe    | Lys   | Leu | Pro        | Gln | Leu | Gln | His | Leu        | Glu | Leu | Asn   | Arg | Asn        |
|        |        |       |     | 155        |     |     |     |     | 160        |     |     |       |     | 165        |
| Lys    | Ile    | Lys   | Asn | Val<br>170 | Asp | Gly | Leu | Thr | Phe<br>175 | Gln | Gly | Leu   | Gly | Ala<br>180 |
| Leu    | Lys    | Ser   | Leu | Lys<br>185 | Met | Gln | Arg | Asn | Gly<br>190 | Val | Thr | Lys   | Leu | Met<br>195 |
| Asp    | Gly    | Ala   | Phe | Trp<br>200 | Gly | Leu | Ser | Asn | Met<br>205 | Glu | Ile | Leu   | Gln | Leu<br>210 |
| Asp    | His    | Asn   | Asn | Leu<br>215 |     | Glu | Ile | Thr | Lys<br>220 | Gly | Trp | Leu   | Tyr | Gly<br>225 |
| Leu    | Leu    | Met   | Leu | Gln<br>230 | Glu | Leu | His | Leu | Ser<br>235 |     | Asn | Ala   | Ile | Asn<br>240 |
| Arg    | Ile    | Ser   | Pro | Asp<br>245 |     | Trp | Glu | Phe | Cys<br>250 |     | Lys | Leu   | Ser | Glu<br>255 |
| Leu    | Asp    | Leu   | Thr | Phe<br>260 |     | His | Leu | Ser | Arg<br>265 |     | Asp | Asp   | Ser | Ser<br>270 |
| Phe    | Leu    | . Gly | Leu | Ser<br>275 |     | Leu | Asn | Thr | Leu<br>280 |     | Ile | : Gly | Asn | Asn<br>285 |
| 7. 200 | . 17-1 | Cor   | TVY | . т1_      | Δ1a | Agn | Cvs | Ala | Phe        | Arc | Glv | Leu   | Ser | Ser        |

|     |     |     |     | 290        |     |     |     |     | 295        |     |     |     |     | 300        |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Leu | Lys | Thr | Leu | Asp<br>305 | Leu | Lys | Asn | Asn | Glu<br>310 | Ile | Ser | Trp | Thr | Ile<br>315 |
| Glu | Asp | Met | Asn | Gly<br>320 | Ala | Phe | Ser | Gly | Leu<br>325 | Asp | Lys | Leu | Arg | Arg<br>330 |
| Leu | Ile | Leu | Gln | Gly<br>335 | Asn | Arg | Ile | Arg | Ser<br>340 | Ile | Thr | Lys | Lys | Ala<br>345 |
| Phe | Thr | Gly | Leu | Asp<br>350 | Ala | Leu | Glu | His | Leu<br>355 | Asp | Leu | Ser | Asp | Asn<br>360 |
| Ala | Ile | Met | Ser | Leu<br>365 | Gln | Gly | Asn | Ala | Phe<br>370 | Ser | Gln | Met | Lys | Lys<br>375 |
| Leu | Gln | Gln | Leu | His<br>380 | Leu | Asn | Thr | Ser | Ser<br>385 | Leu | Leu | Cys | Asp | Cys<br>390 |
| Gln | Leu | Lys | Trp | Leu<br>395 | Pro | Gln | Trp | Val | Ala<br>400 | Glu | Asn | Asn | Phe | Gln<br>405 |
| Ser | Phe | Val | Asn | Ala<br>410 | Ser | Cys | Ala | His | Pro<br>415 | Gln | Leu | Leu | Lys | Gly<br>420 |
| Arg | Ser | Ile | Phe | Ala<br>425 | Val | Ser | Pro | Asp | Gly<br>430 | Phe | Val | Cys | Asp | Asp<br>435 |
| Phe | Pro | Lys | Pro | Gln<br>440 | Ile | Thr | Val | Gln | Pro<br>445 | Glu | Thr | Gln | Ser | Ala<br>450 |
| Ile | Lys | Gly | Ser | Asn<br>455 | Leu | Ser | Phe | Ile | Cys<br>460 | Ser | Ala | Ala | Ser | Ser<br>465 |
| Ser | Asp | Ser | Pro | Met        | Thr | Phe | Ala | Trp | Lys        | Lys | Asp | Asn | Glu | Leu        |
|     |     |     |     | 470        |     |     |     |     | 475        |     |     |     |     | 480        |
| Leu | His | Asp | Ala | Glu<br>485 | Met | Glu | Asn | Tyr | Ala<br>490 | His | Leu | Arg | Ala | Gln<br>495 |
| Gly | Gly | Glu | Val | Met<br>500 | Glu | Туr | Thr | Thr | Ile<br>505 |     | Arg | Leu | Arg | Glu<br>510 |
| Val | Glu | Phe | Ala | Ser<br>515 |     | Gly | Lys | Tyr | Gln<br>520 |     | Val | Ile | Ser | Asn<br>525 |
| His | Phe | Gly | Ser | Ser<br>530 |     | Ser | Val | Lys | Ala<br>535 |     | Leu | Thr | Val | Asn<br>540 |
| Met | Leu | Pro | Ser | Phe<br>545 |     | Lys | Thr | Pro | Met<br>550 |     | Leu | Thr | Ile | Arg<br>555 |

| Ala | Gly | Ala | Met | Ala<br>560 | Arg | Leu | Glu | Cys | Ala<br>565         | Ala | Val | Gly | His | Pro<br>570 |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|--------------------|-----|-----|-----|-----|------------|
| Ala | Pro | Gln | Ile | Ala<br>575 | Trp | Gln | Lys | Asp | Gly<br>580         | Gly | Thr | Asp | Phe | Pro<br>585 |
| Ala | Ala | Arg | Glu | Arg<br>590 | Arg | Met | His | Val | Met<br>595         | Pro | Glu | Asp | Asp | Val<br>600 |
| Phe | Phe | Ile | Val | Asp<br>605 | Val | Lys | Ile | Glu | Asp<br>610         | Ile | Gly | Val | Tyr | Ser<br>615 |
| Cys | Thr | Ala | Gln | Asn<br>620 | Ser | Ala | Gly | Ser | Ile<br>625         | Ser | Ala | Asn | Ala | Thr<br>630 |
| Leu | Thr | Val | Leu | Glu<br>635 | Thr | Pro | Ser | Phe | Leu<br>640         | Arg | Pro | Leu | Leu | Asp<br>645 |
| Arg | Thr | Val | Thr | Lys<br>650 | Gly | Glu | Thr | Ala | Val<br>655         | Leu | Gln | Cys | Ile | Ala<br>660 |
| Gly | Gly | Ser | Pro | Pro<br>665 | Pro | Lys | Leu | Asn | Trp<br>670         | Thr | Lys | Asp | Asp | Ser<br>675 |
| Pro | Leu | Val | Val | Thr<br>680 | Glu | Arg | His | Phe | Phe<br>685         | Ala | Ala | Gly | Asn | Gln<br>690 |
| Leu | Leu | Ile | Ile | Val<br>695 | Asp | Ser | Asp | Val | Ser<br>700         | Asp | Ala | Gly | Lys | Tyr<br>705 |
| Thr | Cys | Glu | Met | Ser<br>710 | Asn | Thr | Leu | Gly | Thr<br>715         | Glu | Arg | Gly | Asn | Val<br>720 |
| Arg | Leu | Ser | Val | Ile<br>725 | Pro | Thr | Pro | Thr | Cys<br>730         | Asp | Ser | Pro | Gln | Met<br>735 |
| Thr | Ala | Pro | Ser | Leu<br>740 | Asp | Asp | Asp | Gly | Trp<br>745         | Ala | Thr | Val | Gly | Val<br>750 |
| Val | Ile | Ile | Ala | Val<br>755 | Val | Cys | Cys | Val | Val<br>760         | Gly | Thr | Ser | Leu | Val<br>765 |
| Trp | Val | Val | Ile | Ile<br>770 | Tyr | His | Thr | Arg | Arg<br><b>7</b> 75 | Arg | Asn | Glu | Asp | Cys<br>780 |
| Ser | Ile | Thr | Asn | Thr<br>785 | Asp | Glu | Thr | Asn | Leu<br>790         | Pro | Ala | Asp | Ile | Pro<br>795 |
| Ser | Tyr | Leu | Ser |            | Gln | Gly | Thr | Leu |                    | Asp | Arg | Gln | Asp |            |

Tyr Val Ser Ser Glu Ser Gly Ser His His Gln Phe Val Thr Ser Ser Gly Ala Gly Phe Phe Leu Pro Gln His Asp Ser Ser Gly Thr 830 Cys His Ile Asp Asn Ser Ser Glu Ala Asp Val Glu Ala Ala Thr Asp Leu Phe Leu Cys Pro Phe Leu Gly Ser Thr Gly Pro Met Tyr 870 860 Leu Lys Gly Asn Val Tyr Gly Ser Asp Pro Phe Glu Thr Tyr His Thr Gly Cys Ser Pro Asp Pro Arg Thr Val Leu Met Asp His Tyr Glu Pro Ser Tyr Ile Lys Lys Glu Cys Tyr Pro Cys Ser His 905 Pro Ser Glu Glu Ser Cys Glu Arg Ser Phe Ser Asn Ile Ser Trp Pro Ser His Val Arg Lys Leu Leu Asn Thr Ser Tyr Ser His Asn 935 Glu Gly Pro Gly Met Lys Asn Leu Cys Leu Asn Lys Ser Ser Leu Asp Phe Ser Ala Asn Pro Glu Pro Ala Ser Val Ala Ser Ser Asn Ser Phe Met Gly Thr Phe Gly Lys Ala Leu Arg Arg Pro His Leu 980 Asp Ala Tyr Ser Ser Phe Gly Gln Pro Ser Asp Cys Gln Pro Arg 1000 Ala Phe Tyr Leu Lys Ala His Ser Ser Pro Asp Leu Asp Ser Gly 1010 1015 Ser Glu Glu Asp Gly Lys Glu Arg Thr Asp Phe Gln Glu Glu Asn 1030 His Ile Cys Thr Phe Lys Gln Thr Leu Glu Asn Tyr Arg Thr Pro 1050 1045 Asn Phe Gln Ser Tyr Asp Leu Asp Thr

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1055

<212> DNA

<213> Homo Sapien

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<211> 640

<212> PRT

<213> Homo Sapien

<400> 292

Met Leu Asn Lys Met Thr Leu His Pro Gln Gln Ile Met Ile Gly
1 5 10 15

Pro Arg Phe Asn Arg Ala Leu Phe Asp Pro Leu Leu Val Val Leu 20 25 30

Leu Ala Leu Gln Leu Leu Val Val Ala Gly Leu Val Arg Ala Gln \$35\$ 40 45

Thr Cys Pro Ser Val Cys Ser Cys Ser Asn Gln Phe Ser Lys Val
50 55 60

Ile Cys Val Arg Lys Asn Leu Arg Glu Val Pro Asp Gly Ile Ser
65 70 75

Thr Asn Thr Arg Leu Leu Asn Leu His Glu Asn Gln Ile Gln Ile 80 85 90

Ile Lys Val Asn Ser Phe Lys His Leu Arg His Leu Glu Ile Leu 95 100 105

Gln Leu Ser Arg Asn His Ile Arg Thr Ile Glu Ile Gly Ala Phe
110 115 120

Asn Gly Leu Ala Asn Leu Asn Thr Leu Glu Leu Phe Asp Asn Arg
125 130 135

Leu Thr Thr Ile Pro Asn Gly Ala Phe Val Tyr Leu Ser Lys Leu 140 145 150

Lys Glu Leu Trp Leu Arg Asn Asn Pro Ile Glu Ser Ile Pro Ser

|     |     |     |     | 155        |     |     |     |     | 160        |     |     |     |     | 165        |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Tyr | Ala | Phe | Asn | Arg<br>170 | Ile | Pro | Ser | Leu | Arg<br>175 | Arg | Leu | Asp | Leu | Gly<br>180 |
| Glu | Leu | Lys | Arg | Leu<br>185 | Ser | Tyr | Ile | Ser | Glu<br>190 | Gly | Ala | Phe | Glu | Gly<br>195 |
| Leu | Ser | Asn | Leu | Arg<br>200 | Tyr | Leu | Asn | Leu | Ala<br>205 | Met | Cys | Asn | Leu | Arg<br>210 |
| Glu | Ile | Pro | Asn | Leu<br>215 | Thr | Pro | Leu | Ile | Lys<br>220 | Leu | Asp | Glu | Leu | Asp<br>225 |
| Leu | Ser | Gly | Asn | His<br>230 | Leu | Ser | Ala | Ile | Arg<br>235 | Pro | Gly | Ser | Phe | Gln<br>240 |
| Gly | Leu | Met | His | Leu<br>245 | Gln | Lys | Leu | Trp | Met<br>250 | Ile | Gln | Ser | Gln | Ile<br>255 |
| Gln | Val | Ile | Glu | Arg<br>260 | Asn | Ala | Phe | Asp | Asn<br>265 | Leu | Gln | Ser | Leu | Val<br>270 |
| Glu | Ile | Asn | Leu | Ala<br>275 | His | Asn | Asn | Leu | Thr<br>280 | Leu | Leu | Pro | His | Asp<br>285 |
| Leu | Phe | Thr | Pro | Leu<br>290 | His | His | Leu | Glu | Arg<br>295 | Ile | His | Leu | His | His<br>300 |
| Asn | Pro | Trp | Asn | Cys<br>305 | Asn | Cys | Asp | Ile | Leu<br>310 | Trp | Leu | Ser | Trp | Trp<br>315 |
| Ile | Lys | Asp | Met | Ala<br>320 | Pro | Ser | Asn | Thr | Ala<br>325 | Cys | Cys | Ala | Arg | Cys<br>330 |
| Asn | Thr | Pro | Pro | Asn<br>335 | Leu | Lys | Gly | Arg | Tyr<br>340 | Ile | Gly | Glu | Leu | Asp<br>345 |
| Gln | Asn | Tyr | Phe | Thr<br>350 | Cys | Tyr | Ala | Pro | Val<br>355 | Ile | Val | Glu | Pro | Pro<br>360 |
| Ala | Asp | Leu | Asn | Val<br>365 | Thr | Glu | Gly | Met | Ala<br>370 | Ala | Glu | Leu | Lys | Cys<br>375 |
| Arg | Ala | Ser | Thr | Ser<br>380 | Leu | Thr | Ser | Val | Ser<br>385 | Trp | Ile | Thr | Pro | Asn<br>390 |
| Gly | Thr | Val | Met | Thr<br>395 | His | Gly | Ala | Tyr | Lys<br>400 | Val | Arg | Ile | Ala | Val<br>405 |
| Leu | Ser | Asp | Gly | Thr        | Leu | Asn | Phe | Thr | Asn        | Val | Thr | Val | Gln | Asp        |

| Thr | Gly | Met | Tyr | Thr<br>425 | Суз | Met | Val | Ser | Asn<br>430 | Ser | Val | Gly | Asn | Thr<br>435 |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Thr | Ala | Ser | Ala | Thr<br>440 | Leu | Asn | Val | Thr | Ala<br>445 | Ala | Thr | Thr | Thr | Pro<br>450 |
| Phe | Ser | Tyr | Phe | Ser<br>455 | Thr | Val | Thr | Val | Glu<br>460 | Thr | Met | Glu | Pro | Ser<br>465 |
| Gln | Asp | Glu | Ala | Arg<br>470 | Thr | Thr | Asp | Asn | Asn<br>475 | Val | Gly | Pro | Thr | Pro<br>480 |
| Val | Val | Asp | Trp | Glu<br>485 | Thr | Thr | Asn | Val | Thr<br>490 | Thr | Ser | Leu | Thr | Pro<br>495 |
| Gln | Ser | Thr | Arg | Ser<br>500 | Thr | Glu | Lys | Thr | Phe<br>505 | Thr | Ile | Pro | Val | Thr<br>510 |
| Asp | Ile | Asn | Ser | Gly<br>515 | Ile | Pro | Gly | Ile | Asp<br>520 | Glu | Val | Met | Lys | Thr<br>525 |
| Thr | Lys | Ile | Ile | Ile<br>530 | Gly | Cys | Phe | Val | Ala<br>535 | Ile | Thr | Leu | Met | Ala<br>540 |
| Ala | Val | Met | Leu | Val<br>545 | Ile | Phe | Tyr | Lys | Met<br>550 | Arg | Lys | Gln | His | His<br>555 |
| Arg | Gln | Asn | His | His<br>560 | Ala | Pro | Thr | Arg | Thr<br>565 | Val | Glu | Ile | Ile | Asn<br>570 |
| Val | Asp | Asp | Glu | Ile<br>575 | Thr | Gly | Asp | Thr | Pro<br>580 | Met | Glu | Ser | His | Leu<br>585 |
| Pro | Met | Pro | Ala | 11e<br>590 | Glu | His | Glu | His | Leu<br>595 | Asn | His | Tyr | Asn | Ser<br>600 |
| Tyr | Lys | Ser | Pro | Phe<br>605 | Asn | His | Thr | Thr | Thr<br>610 | Val | Asn | Thr | Ile | Asn<br>615 |
| Ser | Ile | His | Ser | Ser<br>620 | Val | His | Glu | Pro | Leu<br>625 | Leu | Ile | Arg | Met | Asn<br>630 |
| Ser | Lys | Asp | Asn | Val<br>635 | Gln | Glu | Thr | Gln | Ile<br>640 |     |     |     |     |            |

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<211> 4053

<212> DNA

<213> Homo Sapien

<400> 293

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<211> 1119

<212> PRT

<213> Homo Sapien

<400> 294

Met Ser Ala Pro Ser Leu Arg Ala Arg Ala Ala Gly Leu Gly Leu

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Leu Leu Cys Ala Val Leu Gly Arg Ala Gly Arg Ser Asp Ser Gly 20 25 30

Gly Arg Gly Glu Leu Gly Gln Pro Ser Gly Val Ala Ala Glu Arg

Pro Cys Pro Thr Thr Cys Arg Cys Leu Gly Asp Leu Leu Asp Cys 50 55 60

Ser Arg Lys Arg Leu Ala Arg Leu Pro Glu Pro Leu Pro Ser Trp
65 70 75

Val Ala Arg Leu Asp Leu Ser His Asn Arg Leu Ser Phe Ile Lys

Ala Ser Ser Met Ser His Leu Gln Ser Leu Arg Glu Val Lys Leu
95 100 105

Asn Asn Asn Glu Leu Glu Thr Ile Pro Asn Leu Gly Pro Val Ser 110 115 120

Ala Asn Ile Thr Leu Leu Ser Leu Ala Gly Asn Arg Ile Val Glu 125 130 135

Ile Leu Pro Glu His Leu Lys Glu Phe Gln Ser Leu Glu Thr Leu
140 145 150

Asp Leu Ser Ser Asn Asn Ile Ser Glu Leu Gln Thr Ala Phe Pro 155 160 165

Ala Leu Gln Leu Lys Tyr Leu Tyr Leu Asn Ser Asn Arg Val Thr 170 175 180

Ser Met Glu Pro Gly Tyr Phe Asp Asn Leu Ala Asn Thr Leu Leu 185 190 195

Val Leu Lys Leu Asn Arg Asn Arg Ile Ser Ala Ile Pro Pro Lys 200 205 210

Met Phe Lys Leu Pro Gln Leu Gln His Leu Glu Leu Asn Arg Asn

|     |     |     |     | 215        |     |     |     |     | 220        |     |     |     |     | 225        |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Lys | Ile | Lys | Asn | Val<br>230 | Asp | Gly | Leu | Thr | Phe<br>235 | Gln | Gly | Leu | Gly | Ala<br>240 |
| Leu | Lys | Ser | Leu | Lys<br>245 | Met | Gln | Arg | Asn | Gly<br>250 | Val | Thr | Lys | Leu | Met<br>255 |
| Asp | Gly | Ala | Phe | Trp<br>260 | Gly | Leu | Ser | Asn | Met<br>265 | Glu | Ile | Leu | Gln | Leu<br>270 |
| Asp | His | Asn | Asn | Leu<br>275 | Thr | Glu | Ile | Thr | Lys<br>280 | Gly | Trp | Leu | Tyr | Gly<br>285 |
| Leu | Leu | Met | Leu | Gln<br>290 | Glu | Leu | His | Leu | Ser<br>295 | Gln | Asn | Ala | Ile | Asn<br>300 |
| Arg | Ile | Ser | Pro | Asp<br>305 | Ala | Trp | Glu | Phe | Cys<br>310 | Gln | Lys | Leu | Ser | Glu<br>315 |
| Leu | Asp | Leu | Thr | Phe<br>320 | Asn | His | Leu | Ser | Arg<br>325 | Leu | Asp | Asp | Ser | Ser<br>330 |
| Phe | Leu | Gly | Leu | Ser<br>335 | Leu | Leu | Asn | Thr | Leu<br>340 | His | Ile | Gly | Asn | Asn<br>345 |
| Arg | Val | Ser | Tyr | 11e<br>350 | Ala | Asp | Cys | Ala | Phe<br>355 | Arg | Gly | Leu | Ser | Ser<br>360 |
| Leu | Lys | Thr | Leu | Asp<br>365 | Leu | Lys | Asn | Asn | Glu<br>370 | Ile | Ser | Trp | Thr | Ile<br>375 |
| Glu | Asp | Met | Asn | Gly<br>380 | Ala | Phe | Ser | Gly | Leu<br>385 | Asp | Lys | Leu | Arg | Arg<br>390 |
| Leu | Ile | Leu | Gln | Gly<br>395 | Asn | Arg | Ile | Arg | Ser<br>400 | Ile | Thr | Lys | Lys | Ala<br>405 |
| Phe | Thr | Gly | Leu | Asp<br>410 | Ala | Leu | Glu | His | Leu<br>415 | Asp | Leu | Ser | Asp | Asn<br>420 |
| Ala | Ile | Met | Ser | Leu<br>425 | Gln | Gly | Asn | Ala | Phe<br>430 | Ser | Gln | Met | Lys | Lys<br>435 |
| Leu | Gln | Gln | Leu | His<br>440 | Leu | Asn | Thr | Ser | Ser<br>445 | Leu | Leu | Cys | Asp | Cys<br>450 |
| Gln | Leu | Lys | Trp | Leu<br>455 | Pro | Gln | Trp | Val | Ala<br>460 | Glu | Asn | Asn | Phe | Gln<br>465 |
| Ser | Phe | Val | Asn | Ala        | Ser | Cys | Ala | His | Pro        |     | Leu | Leu | Lys | Gly        |

| Arg | Ser | Ile | Phe | Ala<br>485 | Val | Ser | Pro | Asp | Gly<br>490 | Phe | Val | Cys | Asp | Asp<br>495 |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Phe | Pro | Lys | Pro | Gln<br>500 | Ile | Thr | Val | Gln | Pro<br>505 | Glu | Thr | Gln | Ser | Ala<br>510 |
| Ile | Lys | Gly | Ser | Asn<br>515 | Leu | Ser | Phe | Ile | Cys<br>520 | Ser | Ala | Ala | Ser | Ser<br>525 |
| Ser | Asp | Ser | Pro | Met<br>530 | Thr | Phe | Ala | Trp | Lys<br>535 | Lys | Asp | Asn | Glu | Leu<br>540 |
| Leu | His | Asp | Ala | Glu<br>545 | Met | Glu | Asn | Tyr | Ala<br>550 | His | Leu | Arg | Ala | Gln<br>555 |
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| Arg | Thr | Val | Thr | Lys<br>710 | Gly | Glu | Thr | Ala | Val<br>715 | Leu | Gln | Cys | Ile | Ala<br>720 |
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| Leu | Leu | Ile | Ile | Val<br>755 | Asp | Ser | Asp | Val | Ser<br>760  | Asp | Ala | Gly | Lys | Tyr<br>765        |
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| Glu | Pro | Ser | Tyr | Ile<br>965 | Lys | Lys | Lys | Glu | Cys<br>970  | Tyr | Pro | Cys | Ser | His<br>975        |
| Pro | Ser | Glu | Glu | Ser<br>980 | Cys | Glu | Arg | Ser | Phe<br>985  | Ser | Asn | Ile | Ser | Trp<br>990        |
| Pro | Ser | His | Val | Arg<br>995 | Lys | Leu | Leu |     | Thr<br>1000 | Ser | Tyr | Ser |     | Asn<br>1005       |
| Glu | Gly | Pro | Gly | Met        | Lys | Asn | Leu | Cys | Leu         | Asn | Lys | Ser | Ser | Leu               |

|                           | •                |           | 101     | 0  |      |       |     | 1019            | 5     |     |     | 1020            |
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| Asp A                     | Ala              | Tyr       | Ser Se  |    | Phe  | Gly   | Gln | Pro Se          |       | Cys | Gln | Pro Arg         |
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| Ser (                     | 3lu              | Glu       | Asp G]  |    | Lys  | Glu   | Arg | Thr As          | o Phe | Gln | Glu | Glu Ası<br>109  |
| His 1                     | Ile              | Cys       | Thr Ph  |    | Lys  | Gln   | Thr | Leu Gl          |       | Tyr | Arg | Thr Pro         |
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Asn Thr Tyr Gly Ser Tyr Lys Cys Tyr Cys Leu Asn Gly Tyr Met 110 115 120

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| Ala | Ser   | Thr   | Pro   | Pro<br>365 | Gly | Gly   | Ile   | Thr   | Val<br>370 | Asp | Asn   | Arg | Val | Gln<br>375      |
| Thr | Asp   | Pro   | Gln   | Lys<br>380 |     | Arg   | Gly   | Asp   | Val<br>385 |     | Ser   | Val | Leu | Val<br>390      |
| His | Ser   | Суя   | . Asn | Phe<br>395 |     | His   | Gly   | Leu   | Cys<br>400 |     | Trp   | Ile | Arg | Glu<br>405      |
| Lys | Asp   | ) Asr | n Asp | Leu<br>410 |     | rrp.  | Glu   | Pro   | 11e        |     | Asp   | Pro | Ala | Gl <sub>y</sub> |
| Glv | , Glr | ı Tvi | . Leu |            |     | l Ser | : Ala | ı Ala |            |     | . Pro | Gly | Gly | Ļys             |

|                |       |          |       | 425        |       |       |       |      | 430        |      |     |      |     | 435        |
|----------------|-------|----------|-------|------------|-------|-------|-------|------|------------|------|-----|------|-----|------------|
| Ala            | Ala   | Arg      | Leu   | Val<br>440 | Leu   | Pro   | Leu   | Gly  | Arg<br>445 | Leu  | Met | His  | Ser | Gly<br>450 |
| Asp            | Leu   | Cys      | Leu   | Ser<br>455 | Phe   | Arg   | His   | Lys  | Val<br>460 | Thr  | Gly | Leu  | His | Ser<br>465 |
| Gly            | Thr   | Leu      | Gln   | Val<br>470 | Phe   | Val   | Arg   | Lys  | His<br>475 | Gly  | Ala | His  | Gly | Ala<br>480 |
| Ala            | Leu   | Trp      | Gly   | Arg<br>485 | Asn   | Gly   | Gly   | His  | Gly<br>490 | Trp  | Arg | Gln  | Thr | Gln<br>495 |
| Ile            | Thr   | Leu      | Arg   | Gly<br>500 | Ala   | Asp   | Ile   | Lys  | Ser<br>505 | Glu  | Ser | Gln  | Arg |            |
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| .220.          |       |          |       |            |       |       |       |      |            |      |     |      |     |            |
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| gate           |       |          | gete  | aaqt       | qc c  | ctg : | 24    |      |            |      |     |      |     |            |
| J              |       |          | ,     | J .        | _     | J     |       |      |            |      |     |      |     |            |
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| 005            | 5400  | 990      | 55    |            | -5 -  |       |       |      |            |      |     |      |     |            |
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| <400:          | _ วา  | Ω        |       |            |       |       |       |      |            |      |     |      |     |            |
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| 3              |       | نر       | J     | 5-5        | د     | 2     | _     | ~    |            |      |     | _    |     |            |
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<211> 450

<212> PRT

<213> Homo Sapien

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Gly Val Arg Gly Gln Ala Leu Tyr Leu Pro Val His Tyr Gly Phe

His Thr Pro Ala Ser Asp Ile Gln Ile Ile Trp Leu Phe Glu Arg
50 55 60

| Pro | His | Thr | Met   | Pro<br>65  | Lys | Tyr | Leu | Leu | Gly<br>70  | Ser | Val | Asn | Lys | Ser<br>75  |
|-----|-----|-----|-------|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Val | Val | Pro | Asp   | Leu<br>80  | Glu | Tyr | Gln | His | Lys<br>85  | Phe | Thr | Met | Met | Pro<br>90  |
| Pro | Asn | Ala | Ser   | Leu<br>95  | Leu | Ile | Asn | Pro | Leu<br>100 | Gln | Phe | Pro | Asp | Glu<br>105 |
| Gly | Asn | Tyr | Ile   | Val<br>110 | Lys | Val | Asn | Ile | Gln<br>115 | Gly | Asn | Gly | Thr | Leu<br>120 |
| Ser | Ala | Ser | Gln   | Lys<br>125 | Ile | Gln | Val | Thr | Val<br>130 | Asp | Asp | Pro | Val | Thr<br>135 |
| Lys | Pro | Val | Val   | Gln<br>140 | Ile | His | Pro | Pro | Ser<br>145 | Gly | Ala | Val | Glu | Tyr<br>150 |
| Val | Gly | Asn | Met   | Thr<br>155 | Leu | Thr | Cys | His | Val<br>160 | Glu | Gly | Gly | Thr | Arg<br>165 |
| Leu | Ala | Tyr | Gln   | Trp<br>170 | Leu | Lys | Asn | Gly | Arg<br>175 | Pro | Val | His | Thr | Ser<br>180 |
| Ser | Thr | Tyr | Ser   | Phe<br>185 | Ser | Pro | Gln | Asn | Asn<br>190 | Thr | Leu | His | Ile | Ala<br>195 |
| Pro | Val | Thr | Lys   | Glu<br>200 | Asp | Ile | Gly | Asn | Tyr<br>205 | Ser | Cys | Leu | Val | Arg<br>210 |
| Asn | Pro | Val | Ser   | Glu<br>215 | Met | Glu | Ser | Asp | Ile<br>220 | Ile | Met | Pro | Ile | Ile<br>225 |
| Tyr | Tyr | Gly | Pro   | Tyr<br>230 | Gly | Leu | Gln | Val | Asn<br>235 | Ser | Asp | Lys | Gly | Leu<br>240 |
| Lys | Val | Gly | Glu   | Val<br>245 | Phe | Thr | Val | Asp | Leu<br>250 | Gly | Glu | Ala | Ile | Leu<br>255 |
| Phe | Asp | Суѕ | Ser   | Ala<br>260 |     | Ser | His | Pro | Pro<br>265 | Asn | Thr | Tyr | Ser | Trp<br>270 |
| Ile | Arg | Arg | Thr   | Asp<br>275 | Asn | Thr | Thr | Tyr | Ile<br>280 |     | Lys | His | Gly | Pro<br>285 |
| Arg | Leu | Glu | Val   | Ala<br>290 |     | Glu | Lys | Val | Ala<br>295 |     | Lys | Thr | Met | Asp<br>300 |
| Tyr | Val | Cys | Cys   | Ala<br>305 |     | Asn | Asn | Ile | Thr<br>310 |     | Arg | Gln | Asp | Glu<br>315 |
| Thr | His | Phe | . Thr | Val        | Ile | Ile | Thr | Ser | Val        | Gly | Leu | Glu | Lys | Leu        |

|                                  |              |          |           | 320        |       |       |     |      | 325        |     |     |     |     | 330        |
|----------------------------------|--------------|----------|-----------|------------|-------|-------|-----|------|------------|-----|-----|-----|-----|------------|
| Ala                              | Gln          | Lys      | Gly       | Lys<br>335 | Ser   | Leu   | Ser | Pro  | Leu<br>340 | Ala | Ser | Ile | Thr | Gly<br>345 |
| Ile                              | Ser          | Leu      | Phe       | Leu<br>350 | Ile   | Ile   | Ser | Met  | Cys<br>355 | Leu | Leu | Phe | Leu | Trp<br>360 |
| Lys                              | Lys          | Tyr      | Gln       | Pro<br>365 | Tyr   | Lys   | Val | Ile  | Lys<br>370 | Gln | Lys | Leu | Glu | Gly<br>375 |
| Arg                              | Pro          | Glu      | Thr       | Glu<br>380 | Tyr   | Arg   | Lys | Ala  | Gln<br>385 | Thr | Phe | Ser | Gly | His<br>390 |
| Glu                              | Asp          | Ala      | Leu       | Asp<br>395 | Asp   | Phe   | Gly | Ile  | Tyr<br>400 | Glu | Phe | Val | Ala | Phe<br>405 |
| Pro                              | Asp          | Val      | Ser       | Gly<br>410 | Val   | Ser   | Arg | Ile  | Pro<br>415 | Ser | Arg | Ser | Val | Pro<br>420 |
| Ala                              | Ser          | Asp      | Cys       | Val<br>425 | Ser   | Gly   | Gln | Asp  | Leu<br>430 | His | Ser | Thr | Val | Tyr<br>435 |
| Glu                              | Val          | Ile      | Gln       | His<br>440 | Ile   | Pro   | Ala | Gln  | Gln<br>445 | Gln | Asp | His | Pro | Glu<br>450 |
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| gate                             | cctg         | tca      | caaa      | gcca       | gt g  | gtgc  | 25  |      |            |     |     |     |     |            |
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| <400<br>cac                      |              |          | gtto      | ctca       | icc c | agg:  | 24  |      |            |     |     |     |     |            |
| <210<br><211<br><212             | > 45<br>> DN | ia<br>Ia | i a i a l | 500        | niena | 10    |     |      |            |     |     |     |     |            |

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<212> PRT

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Val Arg Val Ile Thr Asp Glu Asn Trp Arg Glu Leu Leu Glu Gly
35 40 45

Asp Trp Met Ile Glu Phe Tyr Ala Pro Trp Cys Pro Ala Cys Gln 50 55 60

Asn Leu Gln Pro Glu Trp Glu Ser Phe Ala Glu Trp Gly Glu Asp
65 70 75

Leu Glu Val Asn Ile Ala Lys Val Asp Val Thr Glu Gln Pro Gly 80 85 90

Leu Ser Gly Arg Phe Ile Ile Thr Ala Leu Pro Thr Ile Tyr His

Cys Lys Asp Gly Glu Phe Arg Arg Tyr Gln Gly Pro Arg Thr Lys

Lys Asp Phe Ile Asn Phe Ile Ser Asp Lys Glu Trp Lys Ser Ile 125 130 135

Glu Pro Val Ser Ser Trp Phe Gly Pro Gly Ser Val Leu Met Ser 140 145 150

Ser Met Ser Ala Leu Phe Gln Leu Ser Met Trp Ile Arg Thr Cys 155 160 165

His Asn Tyr Phe Ile Glu Asp Leu Gly Leu Pro Val Trp Gly Ser 170 175 180

Tyr Thr Val Phe Ala Leu Ala Thr Leu Phe Ser Gly Leu Leu Leu 185 190 195

Gly Leu Cys Met Ile Phe Val Ala Asp Cys Leu Cys Pro Ser Lys 200 205 210

Arg Arg Pro Gln Pro Tyr Pro Tyr Pro Ser Lys Lys Leu Leu

225 220 215 Ser Glu Ser Ala Gln Pro Leu Lys Lys Val Glu Glu Glu Glu Glu 230 Ala Asp Glu Glu Asp Val Ser Glu Glu Glu Ala Glu Ser Lys Glu 245 Gly Thr Asn Lys Asp Phe Pro Gln Asn Ala Ile Arg Gln Arg Ser 265 Leu Gly Pro Ser Leu Ala Thr Asp Lys Ser <210> 326 <211> 23 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 326 tgaggtgggc aagcggcgaa atg 23 <210> 327 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 327 tatgtggatc aggacgtgcc 20 <210> 328 <211> 21 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 328 tgcagggttc agtctagatt g 21 <210> 329 <211> 25 <212> DNA <213> Artificial Sequence

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Thr Asp Leu Ile Tyr Ala Glu Lys Glu Leu Val Gln Ser Leu Lys
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Glu Tyr Ile Leu Val Glu Glu Ala Lys Leu Ser Lys Ile Lys Ser
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Trp Ala Asn Lys Met Glu Ala Leu Thr Ser Lys Ser Ala Ala Asp
65 70 75

Ala Glu Gly Tyr Leu Ala His Pro Val Asn Ala Tyr Lys Leu Val 80 85 90

Lys Arg Leu Asn Thr Asp Trp Pro Ala Leu Glu Asp Leu Val Leu 95 100 105

Gln Asp Ser Ala Ala Gly Phe Ile Ala Asn Leu Ser Val Gln Arg 110 115 120

Gln Phe Phe Pro Thr Asp Glu Asp Glu Ile Gly Ala Ala Lys Ala 125 130 135

Leu Met Arg Leu Gln Asp Thr Tyr Arg Leu Asp Pro Gly Thr Ile 140 145 150

Ser Arg Gly Glu Leu Pro Gly Thr Lys Tyr Gln Ala Met Leu Ser 155 160 165

Val Asp Asp Cys Phe Gly Met Gly Arg Ser Ala Tyr Asn Glu Gly

Asp Tyr Tyr His Thr Val Leu Trp Met Glu Gln Val Leu Lys Gln 185 190 195

Leu Asp Ala Gly Glu Glu Ala Thr Thr Thr Lys Ser Gln Val Leu 200 205 210

| Asp | Tyr | Leu | Ser | Tyr<br>215 | Ala | Val | Phe | Gln | Leu<br>220 | Gly | Asp | Leu | His | Arg<br>225 |
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| Ala | Leu | Glu | Leu | Thr<br>230 | Arg | Arg | Leu | Leu | Ser<br>235 | Leu | Asp | Pro | Ser | His<br>240 |
| Glu | Arg | Ala | Gly | Gly<br>245 | Asn | Leu | Arg | Tyr | Phe<br>250 | Glu | Gln | Leu | Leu | Glu<br>255 |
| Glu | Glu | Arg | Glu | Lys<br>260 | Thr | Leu | Thr | Asn | Gln<br>265 | Thr | Glu | Ala | Glu | Leu<br>270 |
| Ala | Thr | Pro | Glu | Gly<br>275 | Ile | Tyr | Glu | Arg | Pro<br>280 | Val | Asp | Tyr | Leu | Pro<br>285 |
| Glu | Arg | Asp | Val | Tyr<br>290 | Glu | Ser | Leu | Cys | Arg<br>295 | Gly | Glu | Gly | Val | Lys<br>300 |
| Leu | Thr | Pro | Arg | Arg<br>305 | Gln | Lys | Arg | Leu | Phe<br>310 | Cys | Arg | Tyr | His | His<br>315 |
| Gly | Asn | Arg | Ala | Pro<br>320 | Gln | Leu | Leu | Ile | Ala<br>325 | Pro | Phe | Lys | Glu | Glu<br>330 |
| Asp | Glu | Trp | Asp | Ser<br>335 | Pro | His | Ile | Val | Arg<br>340 | Tyr | Tyr | Asp | Val | Met<br>345 |
| Ser | Asp | Glu | Glu | Ile<br>350 | Glu | Arg | Ile | Lys | Glu<br>355 | Ile | Ala | Lys | Pro | Lys<br>360 |
| Leu | Ala | Arg | Ala | Thr<br>365 | Val | Arg | Asp | Pro | Lys<br>370 | Thr | Gly | Val | Leu | Thr<br>375 |
| Val | Ala | Ser | Tyr | Arg<br>380 | Val | Ser | Lys | Ser | Ser<br>385 | Trp | Leu | Glu | Glu | Asp<br>390 |
| Asp | Asp | Pro | Val | Val<br>395 | Ala | Arg | Val | Asn | Arg<br>400 | Arg | Met | Gln | His | Ile<br>405 |
| Thr | Gly | Leu | Thr | Val<br>410 |     | Thr | Ala | Glu | Leu<br>415 |     | Gln | Val | Ala | Asn<br>420 |
| Tyr | Gly | Val | Gly | Gly<br>425 | Gln | Tyr | Glu | Pro | His<br>430 | Phe | Asp | Phe | Ser | Arg<br>435 |
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| Val | Phe | Pro | Asp | Leu<br>470 |     | Ala | Ala | Ile | Trp<br>475 |     | Lys | Lys | Gly | Thr<br>480 |

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<211> 772

<212> PRT

<213> Homo Sapien

<400> 339

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Val Gly Glu Arg Gly Gly Pro Gln Asn Pro Asp Ser Arg Ala Arg
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Leu Asp Gln Ser Asp Glu Asp Phe Lys Pro Arg Ile Val Pro Tyr
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Tyr Arg Asp Pro Asn Lys Pro Tyr Lys Lys Val Leu Arg Thr Arg 80 85 90

Tyr Ile Gln Thr Glu Leu Gly Ser Arg Glu Arg Leu Leu Val Ala

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|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
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| Asn | Arg | Thr | Val | Ala<br>125 | His | His | Phe | Pro | Arg<br>130 | Leu | Leu | Tyr | Phe | Thr<br>135 |
| Gly | Gln | Arg | Gly | Ala<br>140 | Arg | Ala | Pro | Ala | Gly<br>145 | Met | Gln | Val | Val | Ser<br>150 |
| His | Gly | Asp | Glu | Arg<br>155 | Pro | Ala | Trp | Leu | Met<br>160 | Ser | Glu | Thr | Leu | Arg<br>165 |
| His | Leu | His | Thr | His<br>170 | Phe | Gly | Ala | Asp | Tyr<br>175 | Asp | Trp | Phe | Phe | Ile<br>180 |
| Met | Gln | Asp | Asp | Thr<br>185 | Tyr | Val | Gln | Ala | Pro<br>190 | Arg | Leu | Ala | Ala | Leu<br>195 |
| Ala | Gly | His | Leu | Ser<br>200 | Ile | Asn | Gln | Asp | Leu<br>205 | Tyr | Leu | Gly | Arg | Ala<br>210 |
| Glu | Glu | Phe | Ile | Gly<br>215 | Ala | Gly | Glu | Gln | Ala<br>220 | Arg | Tyr | Cys | His | Gly<br>225 |
| Gly | Phe | Gly | Tyr | Leu<br>230 | Leu | Ser | Arg | Ser | Leu<br>235 | Leu | Leu | Arg | Leu | Arg<br>240 |
| Pro | His | Leu | Asp | Gly<br>245 | Cys | Arg | Gly | Asp | Ile<br>250 | Leu | Ser | Ala | Arg | Pro<br>255 |
| Asp | Glu | Trp | Leu | Gly<br>260 | Arg | Cys | Leu | Ile | Asp<br>265 | Ser | Leu | Gly | Val | Gly<br>270 |
| Cys | Val | Ser | Gln | His<br>275 | Gln | Gly | Gln | Gln | Tyr<br>280 | Arg | Ser | Phe | Glu | Leu<br>285 |
| Ala | Lys | Asn | Arg | Asp<br>290 | Pro | Glu | Lys | Glu | Gly<br>295 | Ser | Ser | Ala | Phe | Leu<br>300 |
| Ser | Ala | Phe | Ala | Val<br>305 |     | Pro | Val | Ser | Glu<br>310 |     | Thr | Leu | Met | Tyr<br>315 |
| Arg | Leu | His | Lys | Arg<br>320 |     | Ser | Ala | Leu | Glu<br>325 |     | Glu | Arg | Ala | Tyr<br>330 |
| Ser | Glu | Ile | Glu | Gln<br>335 |     | Gln | Ala | Gln | Ile<br>340 |     | Asn | Leu | Thr | Val<br>345 |
| Leu | Thr | Pro | Glu | Gly<br>350 |     | Ala | Gly | Leu | Ser<br>355 |     | Pro | Val | Gly | Leu<br>360 |

| Pro | Ala | Pro | Phe | Thr<br>365 | Pro | His | Ser | Arg | Phe<br>370 | Glu | Val | Leu | Gly | Trp<br>375 |
|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
| Asp | Tyr | Phe | Thr | Glu<br>380 | Gln | His | Thr | Phe | Ser<br>385 | Cys | Ala | Asp | Gly | Ala<br>390 |
| Pro | Lys | Cys | Pro | Leu<br>395 | Gln | Gly | Ala | Ser | Arg<br>400 | Ala | Asp | Val | Gly | Asp<br>405 |
| Ala | Leu | Glu | Thr | Ala<br>410 | Leu | Glu | Gln | Leu | Asn<br>415 | Arg | Arg | Tyr | Gln | Pro<br>420 |
| Arg | Leu | Arg | Phe | Gln<br>425 | Lys | Gln | Arg | Leu | Leu<br>430 | Asn | Gly | Tyr | Arg | Arg<br>435 |
|     |     |     |     | 440        |     |     | Glu |     | 445        |     |     |     |     | 450        |
|     |     |     |     | 455        |     |     | His |     | 460        |     |     |     |     | 465        |
|     |     |     |     | 470        |     |     | Ser |     | 475        |     |     |     |     | 480        |
|     |     |     |     | 485        |     |     | Arg |     | 490        |     |     |     |     | 495        |
|     |     |     |     | 500        |     |     | Ala |     | 505        |     |     |     |     | 510        |
|     |     |     |     | 515        |     |     | Arg |     | 520        |     |     |     |     | 525        |
|     |     |     |     | 530        |     |     |     |     | 535        |     |     |     |     | Asp<br>540 |
|     |     |     |     | 545        |     |     |     |     | 550        |     |     |     |     | Arg<br>555 |
| _   |     |     |     | 560        |     |     |     |     | 565        |     |     |     |     | Ala<br>570 |
|     |     |     |     | 575        |     |     |     |     | 580        |     |     |     |     | Pro<br>585 |
|     |     |     |     | 590        |     |     |     |     | 595        |     |     |     |     | Gly<br>600 |
|     |     |     |     | 605        |     |     |     |     | 610        |     |     |     |     | Trp<br>615 |
| Gln | Ala | Phe | Phe | Pro<br>620 |     | His | Phe | Gln | Glu<br>625 |     | Asn | Pro | Ala | 630        |

|   | Ser | Pro   | GIn | Arg | Ser<br>635 | Pro | Pro | Gly | Pro | Pro<br>640 | GIÀ | Ala | GIÀ | Pro | Asp<br>645 |
|---|-----|-------|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|
|   | Pro | Pro   | Ser | Pro | Pro<br>650 | Gly | Ala | Asp | Pro | Ser<br>655 | Arg | Gly | Ala | Pro | 11e        |
|   | Gly | Gly   | Arg | Phe | Asp<br>665 | Arg | Gln | Ala | Ser | Ala<br>670 | Glu | Gly | Cys | Phe | Туг<br>675 |
|   | Asn | Ala   | Asp | Tyr | Leu<br>680 | Ala | Ala | Arg | Ala | Arg<br>685 | Leu | Ala | Gly | Glu | Leu<br>690 |
| • | Ala | Gly   | Gln | Glu | Glu<br>695 | Glu | Glu | Ala | Leu | Glu<br>700 | Gly | Leu | Glu | Val | Met<br>705 |
|   | Asp | Val   | Phe | Leu | Arg<br>710 | Phe | Ser | Gly | Leu | His<br>715 | Leu | Phe | Arg | Ala | Val<br>720 |
| 1 | Glu | Pro   | Gly | Leu | Val<br>725 | Gln | Lys | Phe | Ser | Leu<br>730 | Arg | Asp | Cys | Ser | Pro<br>735 |
|   | Arg | Leu   | Ser | Glu | Glu<br>740 | Leu | Tyr | His | Arg | Cys<br>745 | Arg | Leu | Ser | Asn | 150        |
| • | Glu | Gly   | Leu | Gly | Gly<br>755 | Arg | Ala | Gln | Leu | Ala<br>760 | Met | Ala | Leu | Phe | Glu<br>765 |
| • | Gln | G] 11 | Gln | Ala | Asn        | Ser | Thr |     |     |            |     |     |     |     |            |

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<211> 1572

<212> DNA

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<sup>&</sup>lt;210> 341

<sup>&</sup>lt;211> 318

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

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| Gly               | His  | Gly | Asn | Arg<br>35  | Met | His | His | His | Glu<br>40  | His | His | His | Leu | Gln<br>45  |
| Ala               | Pro  | Asn | Lys | Glu<br>50  | Asp | Ile | Leu | Lys | Ile<br>55  | Ser | Glu | Asp | Glu | Arg<br>60  |
| Met               | Glu  | Leu | Ser | Lys<br>65  | Ser | Phe | Arg | Val | Tyr<br>70  | Cys | Ile | Ile | Leu | Val<br>75  |
| Lys               | Pro  | Lys | Asp | Val<br>80  | Ser | Leu | Trp | Ala | Ala<br>85  | Val | Lys | Glu | Thr | Trp<br>90  |
| Thr               | Lys  | His | Cys | Asp<br>95  | Lys | Ala | Glu | Phe | Phe<br>100 | Ser | Ser | Glu | Asn | Val<br>105 |
| Lys               | Val  | Phe | Glu | Ser<br>110 | Ile | Asn | Met | Asp | Thr<br>115 | Asn | Asp | Met | Trp | Leu<br>120 |
| Met               | Met  | Arg | Lys | Ala<br>125 | Tyr | Lys | Tyr | Ala | Phe<br>130 |     | Lys | Tyr | Arg | Asp<br>135 |
| Gln               | Tyr  | Asn | Trp | Phe<br>140 | Phe | Leu | Ala | Arg | Pro<br>145 | Thr | Thr | Phe | Ala | Ile<br>150 |
| Ile               | Glu  | Asn | Leu | Lys<br>155 | Tyr | Phe | Leu | Leu | Lys<br>160 | Lys | Asp | Pro | Ser | Gln<br>165 |
| Pro               | Phe  | Tyr | Leu | Gly<br>170 | His | Thr | Ile | Lys | Ser<br>175 | Gly | Asp | Leu | Glu | Tyr<br>180 |
| Val               | Gly  | Met | Glu | Gly<br>185 | Gly | Ile | Val | Leu | Ser<br>190 | Val | Glu | Ser | Met | Lys<br>195 |
| Arg               | Leu  | Asn | Ser | Leu<br>200 | Leu | Asn | Ile | Pro | Glu<br>205 | Lys | Cys | Pro | Glu | Gln<br>210 |
| Gly               | Gly  | Met | Ile | Trp<br>215 | Lys | Ile | Ser | Glu | Asp<br>220 | Lys | Gln | Leu | Ala | Val<br>225 |
| Cys               | Leu  | Lys | Tyr | Ala<br>230 | Gly | Val | Phe | Ala | Glu<br>235 | Asn | Ala | Glu | Asp | Ala<br>240 |
| Asp               | Gly  | Lys | Asp | Val<br>245 | Phe | Asn | Thr | Lys | Ser<br>250 | Val | Gly | Leu | Ser | Ile<br>255 |
| Lys               | Glu  | Ala | Met | Thr        | Tyr | His | Pro | Asn | Gln        | Val | Val | Glu | Gly | Сув        |

260 265 270

Cys Ser Asp Met Ala Val Thr Phe Asn Gly Leu Thr Pro Asn Gln 275 280 285

Met His Val Met Met Tyr Gly Val Tyr Arg Leu Arg Ala Phe Gly 290 295 300

His Ile Phe Asn Asp Ala Leu Val Phe Leu Pro Pro Asn Gly Ser 305 310 315

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ttcaaacctg tgatgagaaa aagtttcagc tacctgagaa tttcacagag 300
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gaactgggaa tattttcaat ccagctgcta cttctttct actgacacca 400
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Ile Leu Phe Leu Ser Ala Cys Phe Ile Thr Arg Cys Val Val Thr 35 40 45

Phe Arg Ile Phe Gln Thr Cys Asp Glu Lys Lys Phe Gln Leu Pro 50 55 60

Glu Asn Phe Thr Glu Leu Ser Cys Tyr Asn Tyr Gly Ser Gly Ser
65 70 75

Val Lys Asn Cys Cys Pro Leu Asn Trp Glu Tyr Phe Gln Ser Ser 80 85 90

Cys Tyr Phe Phe Ser Thr Asp Thr Ile Ser Trp Ala Leu Ser Leu
95 100 105

Lys Asn Cys Ser Ala Met Gly Ala His Leu Val Val Ile Asn Ser 110 115 120

Gln Glu Glu Gln Glu Phe Leu Ser Tyr Lys Lys Pro Lys Met Arg 125 130 135

Glu Phe Phe Ile Gly Leu Ser Asp Gln Val Val Glu Gly Gln Trp 140 145 150

Gln Trp Val Asp Gly Thr Pro Leu Thr Lys Ser Leu Ser Phe Trp
155 160 165

Asp Val Gly Glu Pro Asn Asn Ile Ala Thr Leu Glu Asp Cys Ala 170 175 180

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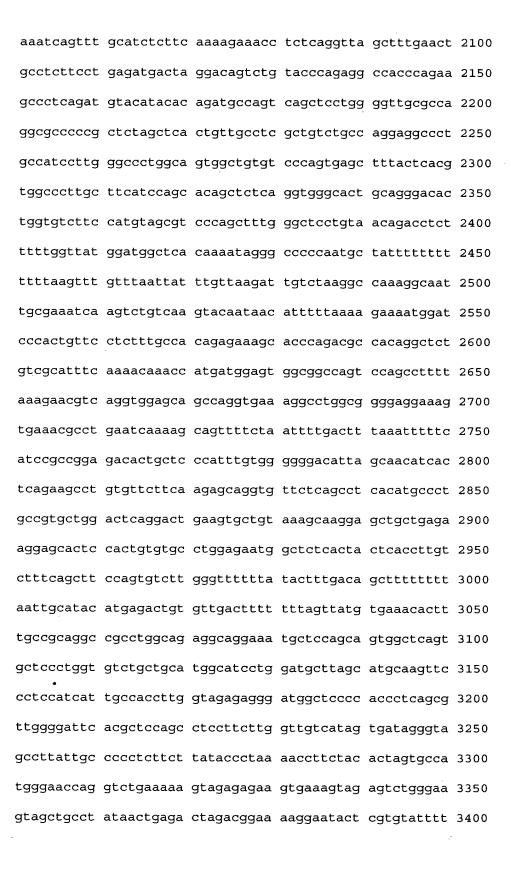
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<212> PRT

<213> Homo Sapien

<400> 423

Met Ala Leu Arg Arg Pro Pro Arg Leu Arg Leu Cys Ala Arg Leu

1 10 15

Pro Asp Phe Phe Leu Leu Leu Phe Arg Gly Cys Leu Ile Gly
20 25 30

Ala Val Asn Leu Lys Ser Ser Asn Arg Thr Pro Val Val Glu Glu 35 40 45

Phe Glu Ser Val Glu Leu Ser Cys Ile Ile Thr Asp Ser Gln Thr
50 55 60

Ser Asp Pro Arg Ile Glu Trp Lys Lys Ile Gln Asp Glu Gln Thr
65 70 75

Thr Tyr Val Phe Phe Asp Asn Lys Ile Gln Gly Asp Leu Ala Gly 80 85 90

Arg Ala Glu Ile Leu Gly Lys Thr Ser Leu Lys Ile Trp Asn Val 95 100 105

Thr Arg Arg Asp Ser Ala Leu Tyr Arg Cys Glu Val Val Ala Arg 110 115 120

Asn Asp Arg Lys Glu Ile Asp Glu Ile Val Ile Glu Leu Thr Val 125 130 135

Gln Val Lys Pro Val Thr Pro Val Cys Arg Val Pro Lys Ala Val 140 145 150

Pro Val Gly Lys Met Ala Thr Leu His Cys Gln Glu Ser Glu Gly
155 160 165

His Pro Arg Pro His Tyr Ser Trp Tyr Arg Asn Asp Val Pro Leu 170 175 180

Pro Thr Asp Ser Arg Ala Asn Pro Arg Phe Arg Asn Ser Ser Phe
185 190 195

| _     |     | •   |     |     | <b>a</b> 1 | m\       | <b>03</b> | mi. | T   | 171        | Db a | mb so | 77.7 | wal | 1114 ~     |
|-------|-----|-----|-----|-----|------------|----------|-----------|-----|-----|------------|------|-------|------|-----|------------|
| ŀ     | HIS | Leu | Asn | Ser | 200        | Tnr      | Gly       | Thr | ьeu | 205        | Pne  | ınr   | Ala  | vai | 210        |
| ·     | Ľуs | Asp | Asp | Ser | Gly<br>215 | Gln      | Tyr       | Tyr | Cys | Ile<br>220 | Ala  | Ser   | Asn  | Asp | Ala<br>225 |
|       | 3ly | Ser | Ala | Arg | Cys<br>230 | Glu      | Glu       | Gln | Glu | Met<br>235 | Glu  | Val   | Tyr  | Asp | Leu<br>240 |
| Z.    | Asn | Ile | Gly | Gly | 11e<br>245 | Ile      | Gly       | Gly | Val | Leu<br>250 | Val  | Val   | Leu  | Ala | Val<br>255 |
| ful I | Leu | Ala | Leu | Ile | Thr<br>260 | Leu      | Gly       | Ile | Cys | Cys<br>265 | Ala  | Tyr   | Arg  | Arg | Gly<br>270 |
|       | Гуr | Phe | Ile | Asn | Asn        | Lys<br>\ | Gln       | Asp | Gly | Glu        | Ser  | Tyr   | Lys  | Asn | Pro        |
|       |     |     |     |     | 275        | $\$      |           |     |     | 280        |      |       |      |     | 285        |
|       | 31y | Lys | Pro | Asp | Gly<br>290 | Val      | Asn       | Tyr | Ile | Arg<br>295 | Thr  | Asp   | Glu  | Glu | Gly<br>300 |
|       | Asp | Phe | Arg | His | Lys<br>305 | ser      | Ser       | Phe | Val | Ile<br>310 |      |       |      |     |            |